## SEQUENCE LISTING

<110> Lodes, Michael J. Mohamath, Raodoh Henderson, Robert A. Benson, Darin R. Secrist, Heather

<120> COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF LUNG CANCER

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stttcagraa ccttttgarc atcetettt ttccgtrtcc eggmaargce cytttccckg ggctttgaaa wyagcetsgt tgggttctta aattaccart ecaenwgttg gaatteeceg ggeeceetge eeggktecaa ecaattttgg graaaaceee encanseegt tkggantgen acaaentggn ntttttentt tegtgnteee etngaaee	540 600 660 698
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 gaaaacaaac agattcatct ccggaaacca aaggaaaggg tragtgggtt tttattagcc
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 agotgtatco tagatggtoa atttocagtg gatgaataca cottacgtac gtttotottg
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                                                                           240
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                                                                           449
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gacaaaaatt tocacccaaa tttgtgcagc taaagcctgg agaaaagcct gttccagtgg gacaaaaatt tocacccaaa tttgtgcagc taaagcctgg agaaacctgag gagaaggaga	540 600 660 720 780 840 900 960 1020 1080 1140 1200 1320 1380 1440 1500 1560 1620 1680 1740 1758
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<210> 28 <211> 1333 <212> DNA <213> Homo sapien

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> <210> 29 <211> 813 <212> DNA <213> Homo sapien

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actggcccgg ggtctgggtc cacctggaca tcgctgctcc agtgcatgct ggcgagcgag 180

15	
ccacaggett tggggtgget etectactgg etetttttgg ecaggaagge gacaacatgg tgetgaacet ggtateeceg etggactgt gagggetaet teecagetgg tgacacaggg eagaggetaet teecagetgg eagaggetaet taageaattg taageaggttg eagaggettgg etgteteett etgtgeed aggggetegg etggggetegg etggggetege etggggetege etggggetege etggggetee etgggggetee etgggggggg	240 300 360 420 480 540 600 660 720 780 813
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cagtecaate atagaaaaga tggaaaaaag gacatgtgcc etgtgeeetg aaggeeacga gtggagtcaa atatacttit caccatcagg aaatatagtt gcicatgaaa actgtttgct 180 gtattcatca ggactggtgg agtgtgagac tettgateta egtaatacaa ttagaaactt 240 tgatgtcaaa tctgtaaaga aagagatctg gagaggaaga agattgaaat gctcattctg 300 taacaaagga ggcgccaccg tggggtgtga tttatggttc tgtaagaaga gttaccacta 360 tgtctgtgcc aaaaaggacc aagcaattct tcaagttgat ggaaaccatg gaacttacaa 420 attattttgc ccagaacatt ctccagaaca agaagaggcc actgaaagtg ctgatgaccc 480 aagcatgaag aagaagagag gaaaaaacaa acgcctctca tcaggccctc ctgcacagcc 540 aaaaacgatg aaatgtagta acgccaaaag acatatgaca gaagagcctc atggtcacac 600 agatgcaget gtcaaatete etttettaa gaaatgceag gaageaggae ttettaetga 660 actatttgaa cacatactag aaaatatgga ttcagttcat ggaagacttg tggatgagac 720 tgcctcagag tcggactatg aagggatcga gaccttactg tttgactgtg gattatttaa 780 agacacacta agaaaattcc aagaagtaat caagagtaaa gcttgtgaat gggaagaaag 840 gcaaaggcag atgaagcage agcttgaggc acttgcagac ttacaacaaa gcttgtgctc 900 atttcaagaa aatggggacc tggactgctc aagttctaca tcaggatcct tgctacctcc 960 tgaggaccac cagtaaaagc tgttcctcag gaaaactgga tggggcctcc atgttctcca 1020 aggatcgagg aagtcttcct gcctaccctg cccaccccag tcaagggcag caacaccaga 1080 gctttgctca gccttaaatg gaatcttaga gctttctctt gcttctgcta ctcctacaga 1140 1200 tggcctcatc atggtctcca ctcagtatta ataactccat cagcatagag caaactcaac actgtgcatt gcacactgtt accatgggtt tatgctcact atcatatcac attgccaata 1260 1316 tttagcacac ttaataaatg cttgtcaaaa cccaaaaaaa aaaaaaaaa ctcgag

> <210> 31 <211> 1355 <212> DNA <213> Homo sapien

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ccaaaggta ttggtggcac ttcaaagatc atgaagagca agataaagtc agacctaaty ccaaaaggaa agaagaacca agctctattt ttcagagaca acgtgtggat gctttacttt tagacctcag acaaaaattt ccacccaaat ttgtgcagct aaagcctgga gaaaacctgag ttccagtgga tcaaacaaga aaagaggcag aacctatacc agaaacctgta aaacctgagg agaaggagac cacaaagaat gtacaacaga cagtgagtgc taaaggccc cctgaaaaac ggatgagact tcagtgagta ctggacaaaa gagaagcctg gaagaccct catgctagtt atcatacctc agtactgtgg ctcttgagct ttgaagtact ttattgtaac cttcttattt atcatacctc agtactgtgg ccatggggg tggatcactt gaggtcagaa gttcaagacc gtaggagac ccatggcggg tggatcactt gaggtcagaa gtcaagacc agcetgacca atatggtgaa accccgtctc tactaaaaat accaaaaatta gccgggggtg agctgacca aggttgccct gagctgatta tcatgctgtt gcactccagc ttgggcgaca cgggaggtgg aggttgccct gagctgatta tcatgctgtt gcactccacac gaacgagact ttgtctcaaa aaaagaagaa aagatattat tcccatcatg atttttccaa	180 540 660 720 780 840 900 960 1020 1080 1140 1260 1320 1355
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Val Ser Arg Ile Arg Gly Gly Ala Lys Lys Arg Lys Lys Ser Tyr  10 15 11 15 10 15 11 10 15 11 10 15 11 10 15 11 10 15 11 10 15 10 15 10 15	
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350
Asn Cys Phe Glu Leu Ser Ile Tyr Val Gly Cys Asp Lys Tyr Pro His
Glu Ser Glu Leu Pro Glu Glu Trp Glu Asn Asn Arg Glu Ser Leu Ile
Val Phe Met Glu Gln Val His Arg Gly Ile Lys Gly Ile Val Arg Asp
Leu Gln Gly Lys Gly Ile Ser Asn Ala Val Ile Ser Val Glu Gly Val
 Asn His Asp Ile Arg Thr Ala Ser Asp Gly Asp Tyr Trp Arg Leu Leu
 Asn Pro Gly Glu Tyr Val Val Thr Ala Lys Ala Glu Gly Phe Ile Thr
 Ser Thr Lys Asn Cys Met Val Gly Tyr Asp Met Gly Ala Thr Arg Cys
 Asp Phe Thr Leu Thr Lys Thr Asn Leu Ala Arg Ile Arg Glu Ile Met
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  Leu Arg Gly Arg Lys Arg Arg Gln Arg Gly
                  485
              500
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   Arg Gly Gln Asp Arg Trp Ser Gln Glu Asp Met Leu Thr Leu Leu Glu
    Cys Met Lys Asn Asn Leu Pro Ser Asn Asp Ser Ser Gln Phe Lys Thr
    Thr Gln Thr His Met Asp Arg Glu Lys Val Ala Leu Lys Asp Phe Ser
    Gly Asp Met Cys Lys Leu Lys Trp Val Glu Ile Ser Asn Glu Val Arg
    Lys Phe Arg Thr Leu Thr Glu Leu Ile Leu Asp Thr Gln Glu His Val
           <210> 36
           <211> 129
           <212> PRT
           <213> Homo sapien
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      Lys Lys Ala Val Ala Ile Ala Asp Ala Leu Gly Lys Ile Pro Gln Thr
      Val Leu Trp Arg Tyr Thr Gly Thr Arg Pro Ser Asn Leu Ala Asn Asn
      Thr Ile Leu Val Gln Trp Leu Pro Gln Asn Asp Leu Leu Gly His Pro
```

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Met Thr Arg Ala Phe Ile Thr His Ala Ser Ser His Gly Val Asn Glu
    Ser Ile Cys Asn Gly Val Pro Met Val Met Ile Pro Leu Phe Gly Asp
    Gln Met Asp Asn Ala Lys Arg Arg Glu Thr Lys Gly Ala Gly Val Thr
    Leu Asn Val Leu Glu Met Thr Ser Glu Asp Leu Glu Asp Ala Leu Lys
     Ser
           <210> 37
           <211> 238
            <212> PRT
            <213> Homo sapien
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      Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe
       Tyr Asp Arg Thr Cys Asn Asn Glu Val Val Lys Met Gln Arg Leu Thr
       Leu Glu His Leu Asn Gln Met Val Gly Ile Glu Tyr Ile Leu Leu His
       Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg Lys Gln Gln Arg Gln Ser
        Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr Tyr Ile Ile Ala Gly Val
        Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val Ile Asn Ser Arg Val Leu
        Thr Ala Val His Gly Ile Gln Ser Ala Phe Asp Glu Ala Met Ser Tyr
ļ:dī
        Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp Trp His Phe Lys Asp His
         Glu Glu Gln Asp Lys Val Arg Pro Lys Ala Lys Arg Lys Glu Glu Pro
         Ser Ser Ile Phe Gln Arg Gln Arg Val Asp Ala Leu Leu Leu Asp Leu
         Arg Gln Lys Phe Pro Pro Lys Phe Val Gln Leu Lys Pro Gly Glu Lys
          Pro Val Pro Val Asp Gln Thr Lys Lys Glu Ala Glu Pro Ile Pro Glu
          Thr Val Lys Pro Glu Glu Lys Glu Thr Thr Lys Asn Val Gln Gln Thr
          Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met Arg Leu Gln
           225
                 <210> 38
                 <211> 202
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<212> PRT

<213> Homo sapien

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Lys Gly Ser Glu Gly Glu Asn Pro Leu Thr Val Pro Gly Arg Glu Lys
Glu Gly Met Leu Met Gly Val Lys Pro Gly Glu Asp Ala Ser Gly Pro
Ala Glu Asp Leu Val Arg Arg Ser Glu Lys Asp Thr Ala Ala Val Val
Ser Arg Gln Gly Ser Ser Leu Asn Leu Phe Glu Asp Val Gln Ile Thr
 Glu Pro Glu Ala Glu Pro Glu Ser Lys Ser Glu Pro Arg Pro Pro Ile
 Ser Ser Pro Arg Ala Pro Gln Thr Arg Ala Val Lys Pro Arg Leu His
 Pro Val Lys Pro Met Asn Ala Thr Ala Thr Lys Val Ala Asn Cys Ser
  Leu Gly Thr Ala Thr Ile Ile Gly Glu Asn Leu Asn Asn Glu Val Met
  Met Lys Lys Tyr Ser Pro Ser Asp Pro Ala Phe Ala Tyr Ala Gln Leu
  Thr His Asp Glu Leu Ile Gln Leu Val Leu Lys Gln Lys Glu Thr Ile
  Ser Lys Lys Glu Phe Gln Val Arg Glu Leu Glu Asp Tyr Ile Asp Asn
  Leu Leu Val Arg Val Met Glu Glu Thr Pro Asn Ile Leu Arg Ile Pro
   Thr Gln Val Gly Lys Lys Ala Gly Lys Met
              180
          195
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          <211> 243
          <212> PRT
          <213> Homo sapien
    Val Asn Ala Leu Gly Ile Met Ala Ala Val Asp Ile Arg Asp Asn Leu
    Leu Gly Ile Ser Trp Val Asp Ser Ser Trp Ile Pro Ile Leu Asn Ser
     Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe Tyr Asp
     Arg Thr Cys Asn Asn Glu Val Val Lys Met Gln Arg Leu Thr Leu Glu
     His Leu Asn Gln Met Val Gly Ile Glu Tyr Ile Leu Leu His Ala Gln
     Glu Pro Ile Leu Phe Ile Ile Arg Lys Gln Gln Arg Gln Ser Pro Ala
      Gln Val Ile Pro Leu Ala Asp Tyr Tyr Ile Ile Ala Gly Val Ile Tyr
      Gln Ala Pro Asp Leu Gly Ser Val Ile Asn Ser Arg Val Leu Thr Ala
      Val His Gly Ile Gln Ser Ala Phe Asp Glu Ala Met Ser Tyr Cys Arg
      Tyr His Pro Ser Lys Gly Tyr Trp Trp His Phe Lys Asp His Glu Glu
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Gln Asp Lys Val Arg Pro Lys Ala Lys Arg Lys Glu Glu Pro Ser Ser 170 Ile Phe Gln Arg Gln Arg Val Asp Ala Leu Leu Leu Asp Leu Arg Gln Lys Ile Ser Thr Gln Ile Cys Ala Val Asp Gln Thr Lys Lys Glu Ala 200 Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met 215 230 Arg Leu Gln

> <210> 40 <211> 245

<212> PRT

<213> Homo sapien

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Ser Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val 40

Val Lys Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly 55

Ile Glu Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile

Arg Lys Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp

Tyr Tyr Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser 105

Val Ile Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala 120

Phe Asp Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr 135

Trp Trp His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys 150

Ala Lys Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val 170

Asp Ala Leu Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val 185

Gln Leu Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys 200

Glu Ala Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr 215

Thr Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys

Arg Met Arg Leu Gln

245

<210> 41 <211> 163 <212> PRT <213> Homo sapien

Gly Glu Arg Gln Gly Leu Val Ala Arg Ala Arg Leu Ser Leu Arg Pro <400> 41 10 Ser Ile Pro Glu Leu Ser Glu Arg Thr Ser Arg Pro Cys Arg Ala Ser 25 Pro Ala Ser Leu Pro Ser Gln His Thr Ser Ser Pro Ala Gln Ala Arg 40 Val Arg Asn Leu Ala Gln Ser Thr Phe Pro Leu Ala Ala Gln Glu Thr 55 Pro Gly Arg Ala Pro Ala His Ala Pro Leu Ser Ser Phe Val Pro Gly 70 Val Gly Gly Arg Ser Pro Ala Ser Val Gly Ile Ser Ala Pro Gly Gly 90 Gly Pro Ser Gly Ala Ala Ala Lys Ile Pro Leu Glu Leu Thr Gln Ser 105 Arg Val Gln Lys Ile Trp Val Pro Val Asp His Arg Pro Ser Leu Pro 120 Arg Ser Cys Gly Pro Lys Leu Thr Asn Ser Pro Ala Val Phe Val Met 140 135 Val Gly Leu Pro Arg Pro Gly Gln Asp Leu Leu Leu His Glu Ser Leu 155 150 Leu Ala Ala

> <210> 42 <211> 243 <212> PRT

<213> Homo sapien

Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp Ser Ser Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val Val Lys 40 Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly Ile Glu 55 Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg Lys 75 Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr Tyr 70 90 Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val Ile 85 105 Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala Phe Asp 120 Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp Trp 135 His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys Ala Lys 155 150 Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val Asp Ala

				165					170	Dro	T.VS	Phe	Val	175 Gln	Leu
Leu	Leu	Leu	Asp	Leu	Arg	Gln	Lys	Phe	Pro	PIO	цуо		190	Gln	
		_	180	T	Dro	Val	Pro	Val	Asp	Gln	Thr	Lys	Lys	Glu	Ala
Lys	Pro	Gly	Glu	гда	PIO	Val	200		_		_	205	mbr	Thr	Lvs
C111	Pro	Tle	Pro	Glu	Thr	Val	Lys	Pro	Glu	Glu	Lys	GIU	1111	1111	Lys
GIU	210					215	70 10	Tue	Glv	Pro	Pro	Glu	Lys	Arg	Met
Asn	Val	Gln	Gln	Thr	Val 230	Ser	Ala	пуз	017	235				Arg	240
225 Arg	Leu	Gln			230										

<210> 43 <211> 244 <212> PRT

<213> Homo sapien

Ala Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp Ser Ser Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val Val Lys Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly Ile 40 Glu Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg Lys Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr Tyr Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val Ile Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala Phe Asp Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp Trp His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys Ala 135 Lys Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val Asp Ala Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val Gln 185 Leu Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys Glu Ala Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr 200 Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met Arg Leu Gln

> <210> 44 <211> 109 <212> PRT

## <213> Homo sapien

Glu Leu His Phe Ser Glu Phe Thr Ser Ala Val Ala Asp Met Lys Asn Ser Val Ala Asp Arg Asp Asn Ser Pro Ser Ser Cys Ala Gly Leu Phe 25 Ile Ala Ser His Ile Gly Phe Asp Trp Pro Gly Val Trp Val His Leu 40 Asp Ile Ala Ala Pro Val His Ala Gly Glu Arg Ala Thr Gly Phe Gly Val Ala Leu Leu Ala Leu Phe Gly Arg Ala Ser Glu Asp Pro Leu 55 Leu Asn Leu Val Ser Pro Leu Asp Cys Glu Val Asp Ala Gln Glu Gly 90 85 Asp Asn Met Gly Arg Asp Ser Lys Arg Arg Arg Leu Val 105

> <210> 45 <211> 324 <212> PRT

<213> Homo sapien

Arg Arg Pro Val Met Ala Gln Glu Thr Ala Pro Pro Cys Gly Pro Val Ser Arg Gly Asp Ser Pro Ile Ile Glu Lys Met Glu Lys Arg Thr Cys

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Ala Leu Cys Pro Glu Gly His Glu Trp Ser Gln Ile Tyr Phe Ser Pro

Ser Gly Asn Ile Val Ala His Glu Asn Cys Leu Leu Tyr Ser Ser Gly Leu Val Glu Cys Glu Thr Leu Asp Leu Arg Asn Thr Ile Arg Asn Phe

Asp Val Lys Ser Val Lys Lys Glu Ile Trp Arg Gly Arg Arg Leu Lys

Cys Ser Phe Cys Asn Lys Gly Gly Ala Thr Val Gly Cys Asp Leu Trp 105

Phe Cys Lys Lys Ser Tyr His Tyr Val Cys Ala Lys Lys Asp Gln Ala 120

Ile Leu Gln Val Asp Gly Asn His Gly Thr Tyr Lys Leu Phe Cys Pro 135

Glu His Ser Pro Glu Gln Glu Glu Ala Thr Glu Ser Ala Asp Asp Pro

Ser Met Lys Lys Lys Arg Gly Lys Asn Lys Arg Leu Ser Ser Gly Pro

Pro Ala Gln Pro Lys Thr Met Lys Cys Ser Asn Ala Lys Arg His Met 185

Thr Glu Glu Pro His Gly His Thr Asp Ala Ala Val Lys Ser Pro Phe 200

Leu Lys Lys Cys Gln Glu Ala Gly Leu Leu Thr Glu Leu Phe Glu His 215

Ile Leu Glu Asn Met Asp Ser Val His Gly Arg Leu Val Asp Glu Thr 235 230

Ala Ser Glu Ser Asp Tyr Glu Gly Ile Glu Thr Leu Leu Phe Asp Cys 250 Gly Leu Phe Lys Asp Thr Leu Arg Lys Phe Gln Glu Val Ile Lys Ser 265 Lys Ala Cys Glu Trp Glu Glu Arg Gln Arg Gln Met Lys Gln Gln Leu 280 Glu Ala Leu Ala Asp Leu Gln Gln Ser Leu Cys Ser Phe Gln Glu Asn 295 Gly Asp Leu Asp Cys Ser Ser Ser Thr Ser Gly Ser Leu Leu Pro Pro 310 Glu Asp His Gln

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Lys Ile Gly Gly Val Thr Glu Arg Met Pro Thr Pro Val Ile Lys Ala
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Leu Asp Pro Lys Ile Ala Asn Ala Ile Met Lys Ala Ala Asp Glu Val
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His Ash Ser Gly Asp Lys Ser Asp 11e Gln Asp Leu Leu Glu Ser Val 50	Ala Gl	n Le	eu G	Glu .	Asn	Glu	Lys	Gln 40	Lys	Val	Ala	GIU	ьец 45	ıyı	261	
Arg Leu Asp Lys Glu Lys Ala Glu Thr Leu Ala Ser Ser Leu Glu Glu Glu Asp Glu Tyr Arg Asn Asp Ala Asn Arg Leu Glu Ala Lys Lys 100 100 105 110 110	His As	n Se	er (	Gly	Asp	Lys	Ser	Asp	Ile	Gln	Asp	Leu 60	Leu	Glu	Ser	Val
Asp Leu Ala His Thr Arg Asn Asp Ala Asn Arg Leu Gln Asp Ala IIe 90 85  Ala Lys Val Glu Asp Glu Tyr Arg Ala Phe Gln Glu Glu Ala Lys Lys 100 105 110 105 110 125  Asp Glu Lys Glu Thr Glu Arg Ser Asp Met Lys Glu Thr IIe Phe Glu 130 135  Leu Glu Asp Glu Val Glu Gln His Arg Ala Val Lys Leu His Asp Asn 155 164  Leu Glu Asp Glu Val Glu Gln His Arg Ala Val Lys Leu His Asp Asn 155 175  Leu IIe IIe Ser Asp Leu Glu Arg Glu Thr Leu His Arg Arg Leu Arg 166  Lys His Asp Met Glu Arg Glu IIe Lys Thr Leu His Arg Arg Leu Arg 180  Glu Glu Ser Ala Glu Trp Arg Gln Phe Gln Ala Asp Leu Gln Thr Ala 205  Val Val IIe Ala Asn Asp IIe Lys Ser Glu Ala Gln Glu Glu IIe Gly 210  Asp Leu Lys Arg Arg Leu His Glu Ala Gln Glu Lys Asn Glu Lys Leu 225  Thr Lys Glu Leu Glu Glu Glu IIe Lys Ser Arg Lys Gln Glu Glu Glu Arg 240  255  Gly Gly Tyr	50 Arg Le	u As	sp 1	Ĺуs	Glu	Lys	Ala	Glu	Thr	Leu	Ala 75	Ser	Ser	Leu	Gln	Glu 80
Ala Lys Val Glu Asp Glu Tyr Arg Ala Phe Gln Glu Glu Ala Lys Lys 100 100 105 110 100 115 120 125 125 125 126 125 126 125 126 126 125 126 126 126 126 126 126 126 126 126 126	65 Asp Le	eu Al	la 1	His	Thr	Arg	Asn	Asp	Ala	Asn	Arg	Leu	Gln	Asp	Ala 95	Ile
Gln Ile Glu Asp Leu Asn Met Thr Leu Glu Lys Leu Arg Ser Asp Leu 115 120 125  Asp Glu Lys Glu Thr Glu Arg Ser Asp Met Lys Glu Thr Ile Phe Glu 130 130 140  Leu Glu Asp Glu Val Glu Gln His Arg Ala Val Lys Leu His Asp Asn 150 150  Leu Ile Ile Ser Asp Leu Glu Asn Thr Val Lys Lys Leu Gln Asp Gln 165  Leu Ile Ile Ser Asp Leu Glu Asn Thr Val Lys Lys Leu Gln Asp Gln 170 175  Lys His Asp Met Glu Arg Glu Ile Lys Thr Leu His Arg Arg Leu Arg 190  Glu Glu Ser Ala Glu Trp Arg Gln Phe Gln Ala Asp Leu Gln Thr Ala 195  Val Val Ile Ala Asn Asp Ile Lys Ser Glu Ala Gln Glu Glu Ile Gly 210 220  Asp Leu Lys Arg Arg Leu His Glu Ala Gln Glu Lys Asn Glu Lys Leu 230 235  Thr Lys Glu Leu Glu Glu Glu Ile Lys Ser Arg Lys Gln Glu Glu Glu Arg 240  245  Gly Gly Tyr <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> <pre> Clo Ser Ala Glu Fre Phe Ser Lys Asn His Ala Ala Pre Phe Ser Lys Val Leu 10 15  Thr Phe Tyr Arg Lys Glu Pre Phe Thr Leu Glu Ala Tyr Tyr Ser Ser 10 15  Thr Phe Tyr Arg Lys Glu Pre Phe Thr Leu Glu Ala Tyr Tyr Ser Ser 20  Pro Gln Asp Leu Pre Tyr Pre Asp Pre Ala Ile Ala Gln Phe Ser Val 40  35  Gln Lys Val Thr Pre Gln Ser Asp Gly Ser Ser Ser Lys Val Lys Val 50  Lys Val Arg Val Asn Val His Gly Ile Phe Ser Val Ser Ser Ala Ser 75  Leu Val Glu Val His Lys Ser Glu Glu Asn Glu Glu Pre Met Glu Thr 100  Pre His Val Glu Glu Glu Gln Gln Gln Gln Thr Pre Gly Arg 125</pre></pre></pre>	Ala Ly	ys Va	al	Glu	85 Asp	Glu	Tyr	Arg	Ala	Phe	Gln	Glu	Glu	Ala 110	Lys	Lys
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Leu Glu Asp Glu Val Glu Gln His Arg Ala Val Lys Leu His Asp Asn 150	Asp G	1 lu L	15 ys	Glu	Thr	Glu	Arg	120 Ser	Ası	o Met	Lys	Glu	Thr	Ile	Phe	Glu
Leu Ile Ile Ser Asp Leu Glu Asn Thr Val Lys Lys Leu Gln Asp Gln  170  185  Lys His Asp Met Glu Arg Glu Ile  185  Lys His Asp Met Glu Arg Glu Ile  185  Lys Thr Leu His Arg Arg Leu Arg  190  Glu Glu Ser Ala Glu Trp Arg Gln Phe Gln Ala Asp Leu Gln Thr Ala  200  Val Val Ile Ala Asn Asp Ile Lys Ser Glu Ala Gln Glu Glu Ile Gly  210  Asp Leu Lys Arg Arg Leu His Glu Ala Gln Glu Lys Asn Glu Lys Leu  230  230  230  231  Thr Lys Glu Leu Glu Glu Ile Lys Ser Arg Lys Gln Glu Glu Arg  245  Gly Gly Tyr	Leu G	30 111 A	asp	Glu	Val	Glu	135 Gln	His	: Ar	g Ala	a Val	Lys	Leu	His	a Asp	Asn 160
Lys His Asp Met Glu Arg Glu Ile Lys Thr Leu His Arg Arg Leu Arg  180  Glu Glu Ser Ala Glu Trp Arg Gln Phe Gln Ala Asp Leu Gln Thr Ala  195  Val Val Ile Ala Asn Asp Ile Lys Ser Glu Ala Gln Glu Glu Ile Gly  210  Asp Leu Lys Arg Arg Leu His Glu Ala Gln Glu Lys Asn Glu Lys Leu  225  Thr Lys Glu Leu Glu Glu Ile Lys Ser Arg Lys Gln Glu Glu Arg  240  Clo Ser  Clo Ser	145	1e T	·le	Ser	Asp	150 Leu	Glu	. Asr	ı Th	r Va	L Lys	o s Lys	Lev	ı Glr	Asp	Gln
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225 Thr Lys Glu Leu Glu Glu Ile Lys Ser Arg Lys Gln Glu Glu Glu Arg 245  Gly Gly Tyr	Val V	7al : 210	Ile	Ala	ASI	n Asj	21	5 5 Б	71	. C1	n G1	22)	o s Asi	n Gl	u Ly	s Leu
Gly Gly Tyr    C210	Asp I 225	Leu :	Lys	Arg	y Aro	g Le <sup>1</sup> 23	и ні. 0	s GT	u Al	.a G1	23	5 (c. G.)	n Gl	n Gl	u Gl	240 u Arg
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Gln Lys Val Thr Pro Gln Ser Asp Gly Ser Ser Lys Val Lys Val 50 55 56 Lys Val Arg Val Asn Val His Gly Ile Phe Ser Val Ser Ser Ala Ser 67 68 69 60 70 75 80 61 62 63 64 65 70 65 65 65 70 65 65 60 70 75 80 65 70 75 80 65 80 65 80 60 70 75 80 80 60 70 75 80 80 61 61 61 61 61 61 61 61 61 61 61 61 61	Thr	Phe	Ту	r Ar 20	) g L	ys G.	Lu Pi	ro P	2	25 25	la T	 1 A	la G	- 3 ln P	0 he S	er Val
50 Lys Val Arg Val Asn Val His Gly Ile Phe Ser Val Ser Ser Ala Ser 65 70 Leu Val Glu Val His Lys Ser Glu Glu Asn Glu Glu Pro Met Glu Thr 85 Asp Gln Asn Ala Lys Glu Glu Glu Lys Met Gln Val Asp Gln Glu Glu 100 Pro His Val Glu Glu Gln Gln Gln Gln Thr Pro Gly Arg 120 125	Pro	Gln	As; 35	p Le	u P	ro T	yr P	ro A 4	sp E 0	ro A		Te u	4.	5 vs V	al L	vs Val
65 Leu Val Glu Val His Lys Ser Glu Glu Asn Glu Glu Pro Met Glu Thr 90 85 85 Asp Gln Asn Ala Lys Glu Glu Glu Lys Met Gln Val Asp Gln Glu Glu 100 100 Pro His Val Glu Glu Gln Gln Gln Thr Pro Gly Arg 120 125	Gln	Lys 50	۷a	l Th	nr P	ro G	ln S 5	er A 5	.sp (	TÀ S	er S	er 5	0 61 n	ys v	or A	la Ser
Leu Val Glu Val His Lys Ser Glu Glu Asn Glu Glu Pro Met Glu Ini 85 90 95 Asp Gln Asn Ala Lys Glu Glu Glu Lys Met Gln Val Asp Gln Glu Glu 100 100 Pro His Val Glu Glu Gln Gln Gln Thr Pro Gly Arg	Lys	Val	Ar	g Va	al A	sn V 7	al H O	is G	ly :	Ile F	he S	ser V 75	aı S	er p	ict V	80
Asp Gln Asn Ala Lys Glu Glu Lys Met Gln Val Asp Gln Glu Glu Glo	bo Leu	Val	. G1	u Va	al H	is L	ys S	er (	lu (	Glu <i>I</i>	Asn G 90	Glu G	lu P	ro M	iet G	5
Pro His Val Glu Glu Gln Gln Gln Thr Pro Gly Alg					la L	ys G									in G L10	ııu Giu
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Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln
Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile
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Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln
Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val
Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly
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 Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu
 Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn
 Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr
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                         215
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   Glu Lys Met Gln Val Asp Gln Glu Glu Pro His Val Glu Glu Gln Gln
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   Gln Gln Thr Pro Ala Glu Asn Lys Ala Glu Ser Glu Glu Met Glu Thr
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Ser Gln Ala Gly Ser Lys Asp Lys Lys Met Asp Gln Pro Pro Gln Ala 75 Lys Lys Ala Lys Val Lys Thr Ser Thr Val Asp Leu Pro Ile Glu Asn 90 Gln Leu Leu Trp Gln Ile Asp Arg Glu Met Leu Asn Leu Tyr Ile Glu 110 105 100 Asn Glu Gly Lys Met Ile Met Gln Asp Lys Leu Glu Lys Glu Arg Asn <210> 62 <211> 418 <212> PRT <213> Homo sapien <400> 62 Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro 10 Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val 20 Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr 40 Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln 55 Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile 75 70 Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val 105 100 Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly 120 Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn 140 135 Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr Ser Leu 155 150 Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly Ala Gly 170 165 Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu 190 185 Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn 205 200 Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr 220 215 Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp Ile Ala 235 230 Thr Ser Gly Ile Ser Thr Thr Phe Pro Lys Leu Arg Met Arg Val Arg 250 245 Asn Ile Leu Ile His Asn Asn Tyr Lys Ser Ala Thr His Glu Asn Asp 265 Ile Ala Leu Val Arg Leu Glu Asn Ser Val Thr Phe Thr Lys Asp Ile 285 280 275 His Ser Val Cys Leu Pro Ala Ala Thr Gln Asn Ile Pro Pro Gly Ser 300 295 Thr Ala Tyr Val Thr Gly Trp Gly Ala Gln Glu Tyr Ala Gly His Thr

305 Val Pro Glu Leu	310 Arg Gln Gly	Gln Val Arg	315 Ile Ile Se		320 Val
Cys Asn Ala Pro	325	330 Asn Gly Ala	Ile Leu Se	er Gly Met	Leu
Cys Asn Ala Pro 340 Cys Ala Gly Val	TIS DEL TYPE	345	Ala Cys Gl	350 In Gly Asp	Ser
Cys Ala Gly Val 355	Pro Gin Giy	360	36	65 rn Phe Ile	Val
355 Gly Gly Pro Leu	. Val Gln Glu 375	Asp Ser Arg	380	- Iva Pro	Glv
370 Gly Ile Val Ser	Trp Gly Asp	Gln Cys Gly	Leu Pro As 395	sp Lys Flo	400
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Cus Met Lys Asn Asn Leu Pro Ser Asn Asp Ser Ser Lys Phe Lys Ini	
35 Thr Glu Ser His Met Asp Trp Glu Lys Val Ala Phe Lys Asp Phe Ser 60	
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Gly Asp Met Cys Lys Hed Lys 1-1 75 80 65 70 75 80 65 Lys Phe Arg Thr Leu Thr Glu Leu Ile Leu Asp Ala Gln Glu His Val 95 95 85 86 86 86 86 86 86 86 86 86 86 86 86 86	
Lys Phe Arg Thr Leu Thr Glu Leu 119 90 95 95 85 90 Asp Phe Pro	
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100 105 Lys Lys Pro Leu Thr Pro Tyr Phe Arg Phe Phe Met Glu Lys Arg Ala Lys Lys Pro Leu Thr Pro Tyr Phe Arg Phe Phe Met Glu Lys Arg Ala 125 120 125	
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150 145 150 175 175 175 177 178 179 170 170 170 170 170 170 170 170 170 170	
Ala Arg Phe Arg Glu Asp His Pro Asp Leu Ile Gln Asn Ala Lys Lys 190	
Ser Asp Ile Pro Glu Lys Pro Lys Thr Pro Gln Gln Leu Trp Tyr Thr	
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Glu Glu Ile Met Arg Asp Tyr Ile Gln Lys His Pro Glu Leu Asn Ile 245 250 265 260 260 260 260 270 260 260	
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Gln Lys Glu Lys Asp Ala Tyr His Lys Lys Cys Asp Gln Lys Lys Asp Tyr Glu Val Glu Leu Leu Arg Phe Leu Glu Ser Leu Pro Glu Glu Glu Gln Gln Arg Val Leu Gly Glu Glu Lys Met Leu Asn Ile Asn Lys 360 Lys Gln Ala Thr Ser Pro Ala Ser Lys Lys Pro Ala Gln Glu Gly Gly 375 Lys Gly Gly Ser Glu Lys Pro Lys Arg Pro Val Ser Ala Met Phe Ile Phe Ser Glu Glu Lys Arg Arg Gln Leu Gln Glu Glu Arg Pro Glu Leu 425 Ser Glu Ser Glu Leu Thr Arg Leu Leu Ala Arg Met Trp Asn Asp Leu 440 Ser Glu Lys Lys Lys Ala Lys Tyr Lys Ala Arg Glu Ala Ala Leu Lys Ala Gln Ser Glu Arg Lys Pro Gly Gly Glu Arg Glu Glu Arg Gly Lys 455 Leu Pro Glu Ser Pro Lys Arg Ala Glu Glu Ile Trp Gln Gln Ser Val Ile Gly Asp Tyr Leu Ala Arg Phe Lys Asn Asp Arg Val Lys Ala Leu 505 Lys Ala Met Glu Met Thr Trp Asn Asn Met Glu Lys Lys Glu Lys Leu 520 Met Trp Ile Lys Lys Ala Ala Glu Asp Gln Lys Arg Tyr Glu Arg Glu Leu Ser Glu Met Arg Ala Pro Pro Ala Ala Thr Asn Ser Ser Lys Lys 535 Met Lys Phe Gln Gly Glu Pro Lys Lys Pro Pro Met Asn Gly Tyr Gln Lys Phe Ser Gln Glu Leu Leu Ser Asn Gly Glu Leu Asn His Leu Pro Leu Lys Glu Arg Met Val Glu Ile Gly Ser Arg Trp Gln Arg Ile Ser Gln Ser Gln Lys Glu His Tyr Lys Lys Leu Ala Glu Glu Gln Gln Lys 600 Gln Tyr Lys Val His Leu Asp Leu Trp Val Lys Ser Leu Ser Pro Gln 615 Asp Arg Ala Ala Tyr Lys Glu Tyr Ile Ser Asn Lys Arg Lys Ser Met Thr Lys Leu Arg Gly Pro Asn Pro Lys Ser Ser Arg Thr Thr Leu Gln 665 Ser Lys Ser Glu Ser Glu Glu Asp Asp Glu Glu Asp Glu Asp Glu 680 Asp Glu Asp Glu Glu Glu Asp Asp Glu Asn Gly Asp Ser Ser Glu Asp Gly Gly Asp Ser Ser Glu Ser Ser Ser Glu Asp Glu Ser Glu Asp Gly Asp Glu Asn Glu Glu Asp Asp Glu Asp Glu Asp Asp Glu Asp 710 730 Asp Asp Glu Asp Glu Asp Asn Glu Ser Glu Gly Ser Ser Ser Ser 745 Ser Ser Leu Gly Asp Ser Ser Asp Phe Asp Ser Asn

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Val Asn Glu Asn Glu Gln Leu Met Glu Asp Tyr Glu Lys Leu Ala
50 55 Ser Asp Leu Glu Trp Ile Arg Arg Thr Ile Pro Trp Leu Glu Asn 80
65 70 73 Arg Val Pro Glu Asn Thr Met His Ala Met Gln Gln Lys Leu Glu Asp 90 95
Pho Arg Asp Tyr Arg Arg Leu His Lys Pro Pro Lys Val Gln Glu Lys
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Thr Arg Asp Leu Cys Ser Leu Val Tyr Val Leu Thr Phe Pro Pro Leu  45
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Thr Ser Leu Gly Thr Asp Lys Cys Glu Ala Leu Leu Gly Leu Cys Gln
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Val Arg Gly Gly Leu Pro Pro Phe Ser Glu Pro Ser Ser Leu Val Pro
Trp Pro Pro Gly Arg Ser Leu Pro Lys Ala Val Arg Pro Pro Leu Ser
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Trp Pro Pro Phe Ser Gln Gln Gln Thr Leu Pro Val Met Ser Gly Glu
                                 105
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 Gly
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 Gly Leu Gln Val Thr Leu Gln Gly Thr Thr Glu Ser Phe Ala Gln Lys
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  Ser Gly Pro Pro Lys Ala Pro Thr Val Ala Glu Gly Pro Ser Ser Cys
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60

780

840 900

960

1500

1560

1620

1680

1740

1800

1860

1920

1980

2040 2076

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<213> Homo sapien

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<212> DNA

<213> Homo sapien

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<211> 386 <212> PRT <213> Homo sapien

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Tyr Thr Val Val Asp Glu Ser Ser Val Ala Lys Ile Asp Asp Ala Ala Pro Pro Glu Lys Val Cys Leu Ile Gly Cys Gly Phe Ser Thr Gly Tyr Gly Ala Ala Val Lys Thr Gly Lys Val Lys Pro Gly Ser Thr Cys Val Val Phe Gly Leu Arg Gly Val Gly Leu Ser Val Ile Met Gly Cys Lys Ser Ala Gly Ala Ser Arg Ile Ile Gly Ile Asp Leu Asn Lys Asp Lys Phe Glu Lys Ala Met Ala Val Gly Ala Thr Glu Cys Ile Ser Pro Lys Asp Ser Thr Lys Pro Ile Ser Glu Val Leu Ser Glu Met Thr Gly Asn Asn Val Gly Tyr Thr Phe Glu Val Ile Gly His Leu Glu Thr Met Ile Asp Ala Leu Ala Ser Cys His Met Asn Tyr Gly Thr Ser Val Val Gly Val Pro Pro Ser Ala Lys Met Leu Thr Tyr Asp Pro Met Leu Leu Phe Thr Gly Arg Thr Trp Lys Gly Cys Val Phe Gly Gly Leu Lys Ser Arg Asp Asp Val Pro Lys Leu Val Thr Glu Phe Leu Ala Lys Lys Phe Asp Leu Asp Gln Leu Ile Thr His Val Leu Pro Phe Lys Lys Ile Ser Glu Gly Phe Glu Leu Leu Asn Ser Gly Gln Ser Ile Arg Thr Val Leu 360 375 Thr Phe 385 <210> 82 <211> 418 <212> PRT <213> Homo sapien Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn

						135					140				
Asn S	130 Ser	Gly	Asn	Leu	Glu	Ile	Asn	Pro	Ser	Thr	Glu	Ile	Thr	Ser	Leu 160
145 Thr 2														Ala	
Pro 2			400					ואה							
Ala		105	Gly				200					200			
Ala	~ 4 ^	His				クコち					220				
Ala					っさい	Ser	Asn			233					
				215	Thr		Phe		2.50						
			$\sim \sim \sim$	His			Tyr	_/nn	Ser				2		
Ile	Ala			Arg	Leu	Glu	Asn 280	Ser	Val	Thr	Phe	Thr 285	Lys	Asp	Ile
His		275 Val	Cys	Leu	Pro	Ala 295	Ala	Thr	Gln	Asn	Ile 300	Pro	Pro	Gly	Ser
	290 Ala	Tyr	Val	Thr	Gly	Trp	Gly	Ala	Gln	Glu 315	Tyr	Ala	Gly	His	Thr 320
305 Val	Pro	Glu	ı Lev	Arg	Gln	Gly	Gln	Val	Arg	ılle	Ile	Ser	Asn	Asp 335	Val
Cys	Asn	Ala	a Pro	325 His	s Ser	туг	Asn	Gly 345	Ala	Ile	Leu	Ser	Gly 350	Met	Leu
Cys	Ala	Gly	340 Val	) L Pro	Glr	Gly	Gly 360	v Val	Asp	Ala	Cys	Gln 365	Gly		Ser
	~ - ~		o Lev			-3.71	ı Asp	Sei			200				val
		va.			201	Ası	o Glr				)				Gly 400
385 Val	Туг	Th	r Ar	g Vai	l Thi	r Ala	а Туз	r Lei	ı Ası 41	o Trp	) Ile	Arg	g Glr	Glr 419	Thr
Gly	, Ile	)		10.	_										

<210> 83

<211> 418

<212> PRT

<213> Homo sapien

<400> 83

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Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg	Asn Gln 95
85 Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser	GIY Vai
Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn	Asn Gly
115 Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met	Leu Asn
130 Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr 155 155	Ser Leu
Asn Ser Gly Asn Leu Glu Tie Ash Tio	160
145 150 153 Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly	175
Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly	, Thr Glu )
180  Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu 205	ı Asn Asn
195 Ala His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile 220 220	e Leu Thr
210 215 220 Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp 230 235	p Ile Ala
Ala Ala His Cys Phe Arg Ser Ash Sel Ash 2235	240
225 230 225 Thr Ser Gly Ile Ser Thr Thr Phe Pro Lys Leu Arg Met Arg	g Val Arg
Thr Ser Gly 11e Ser 112 250	ZDD n Asn Asp
245 Asn Ile Leu Ile His Asn Asn Tyr Lys Ser Ala Thr His Gl	0
260 265  Ile Ala Leu Val Arg Leu Glu Asn Ser Val Thr Phe Thr Ly 285 286	o Clu Ser
275  His Ser Val Cys Leu Pro Ala Ala Thr Gln Asn Ile Pro Pr  295  300	O GIY DOL
295 290 295 Clu Ala Gla Gla Tyr Ala Gl	Ly His Thr
295 296 Thr Ala Tyr Val Thr Gly Trp Gly Ala Gln Glu Tyr Ala Gl 310 315	320
305 Val Pro Glu Leu Arg Gln Gly Gln Val Arg Ile Ile Ser As	sn Asp Val
Val Pro Giu neu 1129 - 1 330 325 330	ວວວ ນ Met Leu
325  Cys Asn Ala Pro His Ser Tyr Asn Gly Ala Ile Leu Ser Gl	50
340  Cys Ala Gly Val Pro Gln Gly Gly Val Asp Ala Cys Gln G	ly Asp Ser
Cys Ala Gly Val Pro Gin Gly Gly 132 365	1. 77.
355  Gly Gly Pro Leu Val Gln Glu Asp Ser Arg Arg Leu Trp Pl	he lie vai
375 380 375 375 380 ASD L	vs Pro Gly
375 370 370 Gly Ile Val Ser Trp Gly Asp Gln Cys Gly Leu Pro Asp L 395	400
385 Val Tyr Thr Arg Val Thr Ala Tyr Leu Asp Trp Ile Arg G	In Gln Thr
Val Tyr Thr Arg val IIII Ald 172 410	415
Gly Ile	

<210> 84 <211> 489 <212> DNA <213> Homo sapien

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120 180

cacgtcggga gcagcttatg agagaagaag ctgaacagaa acgtttaaaa actgtacttg agctacagta tgttttggac aaattgggag atgatgaagt gcggactgac ctgaaacaag gtttgaatgg agtgccaata ttgtccgaag aggagttgtc attgttggat gaattctata agctagtaga ccctgaacgg gacatgagct tgaggttgaa tgaacagtat gaacatgcct ccattcacct gtgggacctg ctggaaggga aggaaaaacc tgtatgtgga accacctata aagttctaa	240 300 360 420 480 489
<210> 85 <211> 304 <212> DNA <213> Homo sapien	
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<210> 86 <211> 296 <212> DNA <213> Homo sapien	
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<210> 88 <211> 387 <212> DNA <213> Homo sapien	
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<210> 91 <211> 488 <212> DNA <213> Homo sapien	
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1 5 10 10 10 10 10 10 10 10 10 10 10 10 10	
20 25 Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu 45 40 45	
35  Ser Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln  55  60	
50  Leu Met Arg Glu Glu Ala Glu Gln Lys Arg Leu Lys Thr Val Leu Glu  80  75  70  75  76  77  78  78  78  78  78  78  78  78	
Leu Gln Tyr Val Leu Asp Lys Leu Gly Asp Asp Glu Val Arg III Asp	
Leu Lys Gln Gly Leu Asn Gly Val Pro Ile Leu Ser Glu Glu Leu	
100 For Leu Leu Asp Glu Phe Tyr Lys Leu Val Asp Pro Glu Arg Asp Met	
115 120 125  Ser Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile His Leu Trp  135 140	
130 135 140  130 135 140  Asp Leu Leu Glu Gly Lys Glu Lys Pro Val Cys Gly Thr Thr Tyr Lys  150 150 160	
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Val Leu	

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> <210> 97 <211> 128 <212> PRT <213> Homo sapien

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> <210> 98 <211> 159 <212> PRT <213> Homo sapien

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Glu Lys Ile Ser Gln Pro Val Val Val Val Ala Ile Val Gly Leu Tyr
Arg Thr Gly Lys Ser Tyr Leu Met Asn His Leu Ala Gly Gln Asn His
Gly Phe Pro Leu Gly Ser Thr Val Gln Ser Glu Thr Lys Gly Ile Trp
 Met Trp Cys Val Pro His Pro Ser Lys Pro Asn His Thr Leu Val Leu
 Leu Asp Thr Glu Gly Leu Gly Asp Val Glu Lys Gly Asp Pro Lys Asn
 Asp Ser Trp Ile Phe Ala Leu Ala Val Leu Cys Ser Thr Phe Val
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  Lys Ile Ser Gln Pro Val Val Val Ala Ile Val Gly Leu Tyr Arg
   Thr Gly Lys Ser Tyr Leu Met Asn His Leu Ala Gly Gln Asn His Gly
   Phe Pro Leu Gly Ser Thr Val Gln Ser Glu Thr Lys Gly Ile Trp Met
   Trp Cys Val Pro His Pro Ser Lys Pro Asn His Thr Leu Val Leu Leu
   Asp Thr Glu Gly Leu Gly Asp Val Glu Lys Gly Asp Pro Lys Asn Asp
    Ser Trp Ile Phe Ala Leu Ala Val Leu Leu Cys Ser Thr Phe Val Tyr
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<213> Homo sapien

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20 25 25 25 20 Ile Asn Asp Pro Phe Ile Asp Leu Asn Tyr Met Val Tyr Met Phe Gln 40 45	
35  Tyr Asp Ser Thr His Gly Lys Phe His Gly Thr Val Glu Ala Glu Asn  55  60	
Tyr Asp Ser III III 557 57 60 50 55 60 Gly Lys Leu Val Ile Asn Gly Asn Pro Ile Thr Ile Phe Gln Glu Arg 80 75 80	
Gly Lys Leu Val lie Ash Gly Ash 75 75 80 65 70 65 Ala Gly Ash Ala Gly Ala Glu Tyr Val Val	
65 70 75  Asp Pro Ser Lys Ile Lys Trp Gly Asp Ala Gly Ala Glu Tyr Val Val  85 90 95  85 Web Cly Lys Ala Gly Ala His Leu	
65  Glu Ser Thr Gly Val Phe Thr Thr Met Glu Lys Ala Gly Ala His Leu  105  100  100  100  100  100  100  10	
Gln Gly Gly Ala Lys Arg Val Ile Ile Ser Ala Pro 115	
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Gly Arg Leu Val Thr Arg Ala Ala 110 45 40 35 40 Asp Leu Asp Tyr Met Val Tyr Met	
35 Val Ala Ile Asn Asp Pro Phe Ile Asp Leu Asn Tyr Met Val Tyr Met Val Ala Ile Asn Asp Pro Phe Ile Asp Leu Asn Tyr Met Val Glu Ala	
55 50 50 50 Fig. 10 Fig. 11 Fi	
65 Glu Asn Gly Lys Leu Val Ile Asn Gly Asn Pro Ile Thr Ile Phe Gln 95 90 85	
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acananacaa cadcogugac ggggg	180 240
acttagaca accepyyyoo ososoo a acataataa acceages	300 360
gactectgeg accegeaceg caceceace egggeeegga gggeegeatg cetgtgttte aaccagacee geacetacga eggegaegge tacaagaage gggeegeatg cetgtgttte	500

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l.al.

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 Lys Thr Ser Ser Thr Lys Ser Ala Ile Cys Ala Arg Ala Ile Pro Ala
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 Glu Leu Trp Ser Leu Phe Asp Phe Ala Cys Gln Gly Ser Leu Leu Gly
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 Arg Glu Lys Asp Ala Thr Pro Gly Glu Lys Ala Leu Gly Phe Lys Ile
                             120
 Ser Glu Asn Leu Met Ala Ile Ile Lys Pro Tyr Phe Leu Arg Arg Thr
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 Lys Glu Asp Val Gln Lys Lys Ser Ser Asn Pro Glu Ala Arg Leu
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 Asn Glu Lys Asn Pro Asp Val Asp Ala Ile Cys Glu Met Pro Ser Leu
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 Ser Arg Arg Asn Asp Leu Ile Ile Trp Ile Arg Leu Val Pro Leu Gln
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  Glu Glu Ile Tyr Arg Lys Phe Val Ser Leu Asp His Ile Lys Glu Leu
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                             200
  Leu Met Glu Thr Arg Ser Pro Leu Ala Glu Leu Gly Val Leu Lys Lys
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                          215
  Leu Cys Asp His Pro Arg Leu Leu Ser Ala Arg Ala Cys Cys Leu Leu
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65 70 75 80  65 Tyr Asp Met Arg Leu Arg Arg  Met Glu Tyr Lys Gly Glu Leu Ala Ser Tyr Asp Met Arg Leu Arg Arg  90 95  85 91 The His Val Lys Ser Leu Pro Gly	
Met Glu Tyl Bys 90  85  Lys Leu Asp Leu Phe Ala Asn Val Ile His Val Lys Ser Leu Pro Gly 105  100  Tyr Met Thr Arg His Asn Asn Leu Asp Leu Val Ile Ile Arg Glu Gln 120  Thr	
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100

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Ser Leu His Leu Ser Pro Gln Val Ser Ser Ser Val Val Tyr Gly Arg
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Ser Pro Val Met Ala Phe Glu Ser His Pro His Leu Arg Gly Ser Ser
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Val Ser Ser Ser Leu Pro Ser Ile Pro Gly Gly Lys Pro Ala Tyr Ser
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                              40
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  Cys Ser Asp Arg Thr Ser Pro Tyr Asp His Met Leu Pro Gly Ala Glu
  His Phe Ala Glu Tyr Ala Gly Arg Leu Gly Val Gly Ala Ala Thr His
  Val Val Ile Tyr Asp Ala Ser Asp Gln Gly Leu Tyr Ser Ala Pro Arg
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  Val Trp Trp Met Phe Arg Ala Phe Gly His His Ala Val Ser Leu Leu
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  Asp Gly Gly Leu Arg His Trp Leu
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Arg Ser Glu Thr Ser Val Pro Asp His Val Val Trp Ser Leu Phe Asn
Thr Leu Phe Met Asn Pro Cys Cys Leu Gly Phe Ile Ala Phe Ala Tyr
Ser Val Lys Ser Arg Asp Arg Lys Met Val Gly Asp Val Thr Gly Ala
 Gln Ala Tyr Ala Ser Thr Ala Lys Cys Leu Asn Ile Trp Ala Leu Ile
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  Ser Thr Pro Gly Thr Ala Pro Pro Pro Lys Val Leu Thr Ser Pro Ala
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  Glu Thr Thr His Thr Ser Thr Val Leu Thr Thr Thr Ala Thr Met Thr
  Arg Ala Thr Asn Ser Thr Ala Thr Pro Ser Ser Thr Leu Gly Thr Thr
   Arg Ile Leu Thr Glu Leu Thr Thr Thr Ala Thr Thr Thr Ala Ala Thr
   Gly Ser Thr Ala Thr Leu Ser Ser Thr Pro Gly Thr Thr Trp Ile Leu
   Thr Glu Pro Ser Thr Ile Ala Thr Val Met Val Pro Thr Gly Ser Thr
   Ala Thr Ala Ser Ser Thr Leu Gly Thr Ala His Thr Pro Lys Val Val
   Thr Thr Met Ala Thr Met Pro Thr Ala Thr Ala Ser Thr Val Pro Ser
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- at at analog Cayocacyau	660 720
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Ala Leu Gin Lys Bed 25 20  Ala Lys Lys Lys Val Leu Phe Ala Leu Cys Ser Leu Leu Arg His Phe 45 35  Pro Tyr Ala Gln Arg Gln Phe Leu Lys Leu Gly Gly Leu Gln Val Leu 50  Arg Thr Leu Val Gln Glu Lys Gly Thr Glu Val Leu Ala Val Arg Val 70  Not Phe Ala Glu Glu	
Arg Thr Leu Val Gin Gid 57  65  70  Val Thr Leu Leu Tyr Asp Leu Val Thr Glu Lys Met Phe Ala Glu Glu 95  85  Glu Ala Glu Leu Thr Gln Glu Met Ser Pro Glu Lys Leu Gln Gln Tyr 110  105  105  107  108  109  109  100  100  100  100  100	
Glu Ala Glu Leu III 31 105 116 117 117 118 119 119 119 119 119 119 119 119 119	
130 The Lou Cly Val Leu Leu Thr Thr Cys Arg Asp 160	
Val Leu Gln Thr Leu Gly  155  145  Arg Gln Asp Pro Gln Leu Gly Arg Thr Leu Ala Ser Leu Gln Ala Glu  175  165  Tyr Gln Val Leu Ala Ser Leu Glu Leu Gln Asp Gly Glu Asp Glu Gly  180  180  180  180  180  180  180  18	
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245	255	
Trp Gly Val Ala Thr Ala Ser Leu Ile Pro Asn Asp Gln Leu Leu 170 275 280 280 280 285 285 285 285 285 285 295 295 295 295 295 295 295 295 295 29	250 250 250	ar
Trp Gly Val Ala Thr Ala Ser Leu Ile Pro Asn Asp Gln Leu Leu 170 275 280 280 280 285 285 285 285 285 285 295 295 295 295 295 295 295 295 295 29	245  Win Car Leu Thr Ser Gln Ser Gln Ile Ser Pro Lys Se	21
Trp Gly Val Ala Thr Ala Ser Leu Ile Pro Asn Asp Gln Leu Leu 170 275 280 280 280 285 285 285 285 285 285 295 295 295 295 295 295 295 295 295 29	Ser Gln Lys His Ser neu 1112 265	**
Arg Lys Leu Asn Thr Glu Pro Lys Asp Val Pro Lys Pro Val Ris Glu 290  Pro Val Gly Ser Ser Ser Thr Leu Pro Lys Asp Pro Val Leu Arg Lys 315  Glu Lys Leu Gln Asp Leu Met Thr Gln IIe Gln Gly Thr Cys Asn Phe 325  Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala IIe Pro 340  Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 360  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 375  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu IIe 400  385  Glu Gln Thr Val His Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 435  A 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 440  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 465  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 400  A 455  Asn Val Pro Val Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser Ser Ser Gln Gln Tyr Arg Gly Leu Pro Ser 555  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys Gly Gly Tyr Lys Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 495  Arg Gly Gly Thr Ser Gln Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Asp Ser Arg Ser Met Thr Pro 555  Ang Gly Gly Thr Ser Gln Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Arg Gly Gly Thr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gln Gln Met Arg Val Ala Phe Ser Arg Ser Met Thr Pro Val 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Glo	260 Asn Asp Gln Leu Leu F.	10
Arg Lys Leu Asn Thr Glu Pro Lys Asp Val Pro Lys Pro Val Ris Glu 290  Pro Val Gly Ser Ser Ser Thr Leu Pro Lys Asp Pro Val Leu Arg Lys 315  Glu Lys Leu Gln Asp Leu Met Thr Gln IIe Gln Gly Thr Cys Asn Phe 325  Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala IIe Pro 340  Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 360  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 375  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu IIe 400  385  Glu Gln Thr Val His Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 435  A 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 440  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 465  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 400  A 455  Asn Val Pro Val Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser Ser Ser Gln Gln Tyr Arg Gly Leu Pro Ser 555  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys Gly Gly Tyr Lys Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 495  Arg Gly Gly Thr Ser Gln Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Asp Ser Arg Ser Met Thr Pro 555  Ang Gly Gly Thr Ser Gln Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Arg Gly Gly Thr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gln Gln Met Arg Val Ala Phe Ser Arg Ser Met Thr Pro Val 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val Glu Glo	Trp Gly Val Ala Thr Ala Sel Box 280 285	1
290 Pro Val Gly Ser Ser Ser Thr Leu Pro Lys Asp Pro Val Leu Au 320 310 310 311 310 315 316 317 318 320 321 Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala Ile Pro 340 340 Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 355 Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370 375 Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 370 Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 370 Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 405 Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 405 Cln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 406 Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 407 Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 408 Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 500 Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 505 Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Arg Asp Asn Ser Arg Ala Gly Trp 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Arg Asp Asn Ser Arg Ala Gly Trp 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 506 Ser Asp Ser Gly Asp Ser Gly Gly Pro Arg Ala Ash Ser Arg Ala Gly Trp 506 Ser Asp Ser Gly Asp Ser Gly Gly Pro Arg Ala Ash Ser Arg Ala Gly Trp 507 Ser Ash Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 508 Ser Asp Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 509 Ser Ash Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610 Ser Ash Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 610 Ser Ash Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 610 Ser Ash Leu Ala Pro Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	275 Lys Asp Val Pro Lys Pro Val His G	T11
290 Pro Val Gly Ser Ser Ser Thr Leu Pro Lys Asp Pro Val Leu Au 320 310 310 311 310 315 316 317 318 320 321 Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala Ile Pro 340 340 Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 355 Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370 375 Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 370 Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 370 Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 405 Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 405 Cln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 406 Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 407 Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 408 Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 500 Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 505 Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Arg Asp Asn Ser Arg Ala Gly Trp 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Gly Arg Asp Asn Ser Arg Ala Gly Trp 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 505 Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Tyr 506 Ser Asp Ser Gly Asp Ser Gly Gly Pro Arg Ala Ash Ser Arg Ala Gly Trp 506 Ser Asp Ser Gly Asp Ser Gly Gly Pro Arg Ala Ash Ser Arg Ala Gly Trp 507 Ser Ash Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 508 Ser Asp Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 509 Ser Ash Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610 Ser Ash Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 610 Ser Ash Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 610 Ser Ash Leu Ala Pro Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	Arg Lys Leu Asn Thr Glu Pro Lys 100 300	
Solution   Solution	290 Zys Lou Pro Lys Asp Pro Val Leu Arg L	ys.
Solution   Solution	Pro Val Gly Ser Ser Ser Thr Leu Flo Bys 135	320
Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala The To 340  Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 355  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 400  385  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 415  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 440  435  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460  Asn Val Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser Asp Val Pro Asp Gly Tyr Lys Gly Phe Asp Thr Tyr Asn Gln Leu Pro Ser Tyr Arg 490  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 550  Asn Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Gly Arg Leu Ile Thr Asn Ser Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 550  Asn Ser Gly Asp Ser Gly Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 600  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 600  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Leu Leu Asn Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 655  Leu Leu Asn Asn Leu Gly Glu Thr Pro Pro Ala Ala Cly Tyr Val Phe Leu Gly Arg Phe 1650  As a Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 655	310 Sla Tlo Gla Gla Thr Cys Asn F	Phe
Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala The To 340  Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 355  Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 400  385  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 415  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 440  435  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460  Asn Val Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser Asp Val Pro Asp Gly Tyr Lys Gly Phe Asp Thr Tyr Asn Gln Leu Pro Ser Tyr Arg 490  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 550  Asn Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Gly Arg Leu Ile Thr Asn Ser Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 550  Asn Ser Gly Asp Ser Gly Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Tyr Lys 560  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 600  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 600  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Leu Leu Asn Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 655  Leu Leu Asn Asn Leu Gly Glu Thr Pro Pro Ala Ala Cly Tyr Val Phe Leu Gly Arg Phe 1650  As a Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 655	Clu Ivs Leu Gln Asp Leu Met Thr Gln 11e Gln Gly 200 335	
Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 360 365 365 365 365 360 370 370 370 370 370 390 390 390 400 385 400 390 395 400 385 405 405 405 405 405 405 405 405 405 40	325 Son Son Ser Ala Ile F	?ro
Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys 360 365 365 365 365 360 370 370 370 370 370 390 390 390 400 385 400 390 395 400 385 405 405 405 405 405 405 405 405 405 40	350	
Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 380  390  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 410  Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 435  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460  A50  A55  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 465  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Tyr Arg 465  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 515  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 555  Ser Asp Ser Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 662  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 663  660  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 665  Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	Met Gin Giu Scr van 345	โเซร
Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln 370  Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile 380  390  Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala 410  Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu 420  Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 435  Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460  A50  A55  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 465  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Tyr Arg 465  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 515  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 555  Ser Asp Ser Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 662  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 663  660  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 665  Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Bor 2	2,10
No	Thr Ser Gin Pio Pio Boz 360	Gln
No	355 Car Sar Gln Ser Asp Phe Leu Gln Glu Pro Leu	O 1 1 1
No	Glu Gln Asn Leu Ser Ser Sin 375	т10
Ser   Ser   Pro   Tyr   Ser   Pro   Gly   Tyr   Ash   Gln   Ser   Phe   Thr   Thr   Ala	370 Pro Leu Pro Pro Arg Lys Glu Gln Glu	TTE
Ser   Ser   Pro   Tyr   Ser   Pro   Gly   Tyr   Ash   Gln   Ser   Phe   Thr   Thr   Ala	Val Phe Asn Val Asn Ala Plo Hed 123 395	400
Ser Thr Gln Thr         Pro Pro Gln Gln Cys         Gln Leu Pro Ser IIe His Val Gln 430         430           Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 435         440         445           Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460         455         460           Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 480         465         480           Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 490         495           Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510         500           Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 520         525           Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530         535           Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 560         555           Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 555         560           Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580         580           Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605         605           Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620         605           Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo	385 390 Cly Tyr Asn Gln Ser Phe Thr Thr	Ala
Ser Thr Gln Thr         Pro Pro Gln Gln Cys         Gln Leu Pro Ser IIe His Val Gln 430         430           Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly 435         440         445           Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr 460         455         460           Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser 480         465         480           Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 490         495           Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510         500           Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 520         525           Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530         535           Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 560         555           Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 555         560           Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580         580           Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605         605           Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620         605           Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo	Lys Glu Ser Pro Tyr Ser Pro Gly Tyl Abh 415	
Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly  435  Thr Ile Gln Val Ser Asn Gly  450  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly  465  Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg  485  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser  500  Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr  520  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys  530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Gly Trp  530  Arg Gly Gly Thr Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr  565  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val  580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val  600  590  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr  610  Ser Asn Leu Ala Pro Gly Glu Thr Tyr Val Phe His Met Leu Lys  640  645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe His Met Leu Lys  6670	405 and Low Pro Ser Ile His Val	Glu
Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly  435  Thr Ile Gln Val Ser Asn Gly  450  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly  465  Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg  485  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser  500  Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr  520  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys  530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Gly Trp  530  Arg Gly Gly Thr Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr  565  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val  580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val  600  590  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr  610  Ser Asn Leu Ala Pro Gly Glu Thr Tyr Val Phe His Met Leu Lys  640  645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe His Met Leu Lys  6670	cor Thr Gln Thr Pro Pro Gln Cys Gln Leu 110 301 430	
Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Interval 450 455 466 455 466 467 470 488 475 480 475 480 475 480 470 485 485 485 485 485 485 485 485 485 485	425 420 425 Asp	Gly
Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Interval 450 455 466 455 466 467 470 488 475 480 475 480 475 480 470 485 485 485 485 485 485 485 485 485 485	The Wal His Ser Glu Glu Thr Ala Ala Ash Tyl Mis 245	
Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly 361  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly 361  465  Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 495  485  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510  500  Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 550  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 585  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo 625  Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	440 440 Ala Gln	Thr
Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly 361  Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly 361  465  Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg 495  485  Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510  500  Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 550  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 585  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo 625  Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	433 Ser Asn Gly Ser Leu Ala Phe Tyr Pio Ala Gin	
465       Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyl Arg 495         85       485       490       495       495         Ser Pro Gly Gly Tyr Lys Gly Phe Sp 505       500       505       510       510         Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525       520       525       525         Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530       535       540       540       550         Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asp Ser Arg Ser Arg Ala Gly Trp 550       550       555       560       560         Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 570       575       575       575       575         Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 590       585       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       595       500       595       500       500       500       500	Thr Ile Gin var Scr 1355 460	Ser
465       Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyl Arg 495         85       485       490       495       495         Ser Pro Gly Gly Tyr Lys Gly Phe Sp 505       500       505       510       510         Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525       520       525       525         Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530       535       540       540       550         Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asp Ser Arg Ser Arg Ala Gly Trp 550       550       555       560       560         Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 570       575       575       575       575         Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 590       585       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       595       500       595       500       500       500       500	450 The Bro The Gln Pro Phe Val Asn Ser Arg Gly	780
465       Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyl Arg 495         85       485       490       495       495         Ser Pro Gly Gly Tyr Lys Gly Phe Sp 505       500       505       510       510         Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525       520       525       525         Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530       535       540       540       550         Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asp Ser Arg Ser Arg Ala Gly Trp 550       550       555       560       560         Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 570       575       575       575       575         Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 590       585       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       585       590       590       590       585       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       590       595       500       595       500       500       500       500	Asn Val Phe Pro Arg 110 111 475	700
Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510  1le Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 550  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 565  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 575  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 615  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620  Ser Asn Leu Ala Pro Gly Glu Thr Phe Asp Leu Gly Arg Phe Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	465 Arg Cly Gly Arg Leu Ile Thr Asn Ser Tyl	ALG
Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser 510  1le Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr 525  Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys 530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp 550  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 565  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 575  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 615  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620  Ser Asn Leu Ala Pro Gly Glu Thr Phe Asp Leu Gly Arg Phe Leu Leu Asn Asn Leu Gly Glu Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	Val Arg Gly Cys Thr Arg Gry Ori 490	0
Ser   Asn   Gly   Asn   Tyr   Ser   Gln   Leu   Gln   Phe   Gln   Ala   Arg   Glu   Tyr	485 Cly Phe Asp Thr Tyr Arg Gly Leu Pro	Ser
Ser   Asn   Gly   Asn   Tyr   Ser   Gln   Leu   Gln   Phe   Gln   Ala   Arg   Glu   Tyr	Ser Pro Gly Gly Tyr Lys Gly File 105 510	
Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys  530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp  550  550  560  570  570  570  575  570  575  570  570  575  570  580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val  595  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr  610  527  538  540  540  540  550  550  550  560  570  570  575  575	500 Garage Clar Lou Glar Phe Glar Ala Arg Glu	ı Tyr
Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys  530  Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp  550  550  560  570  570  570  575  570  575  570  570  575  570  580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val  595  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr  610  527  538  540  540  540  550  550  550  560  570  570  575  575	The Ser Ash Gly Ash Tyr Ser Gin Led Gin 2505	
Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly 115  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 600  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 635  Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 655  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	515 San Den Phe Gln Gln Cys Tyl	r Lys
Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly 115  Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe 575  Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val 580  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 600  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 635  Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 655  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	Ser Gly Ala Pro Tyr Ser Gln Arg Asp Ash The 540	
545       Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Intraction       570       575         Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val       580       585       590         Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val       605       605         Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr       620       620         610       615       620       640         Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo G40       625       640         Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe G55       655         Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	535 530 535 Fig. 7 and Ser Arg Ala Gl.	y Trp
545       Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Intraction       570       575         Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val       580       585       590         Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val       605       605         Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr       620       620         610       615       620       640         Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu Glo G40       625       640         Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe G55       655         Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	and Gly Gly Thr Ser Gly Gly Pro Arg Ala Ash Gol 11-9	560
Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr F16 various 580 580 585 590  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605 605 600 605 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620 610 615 625 645 630 635 640 655 655 645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	550 St. Amg Ash Glu Th	r Phe
Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr F16 various 580 580 585 590  Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 605 605 600 605 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 620 610 615 625 645 630 635 640 655 655 645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys 670	345 Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Ash 57	5
Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 595 600 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610 615 620  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 635 640  Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 650 655  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	Ser Asp Ser John 565	o Val
Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 595 600 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610 615 620  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 635 640  Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 650 655  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	Gran Cly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Int II	
Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val 595 600 605  Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr 610 615 620  Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu 635 640  Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 650 655  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	Asn Ser Gly Asp 501 5-1 585 590	e Val
Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg III 620 610 615 625 640 625 620 640 630 640 625 640 625 645 645 650 650 655 645 650 6670	The Asp Pro Ala Ala Thr Ile Leu Pro Val Hi	J VUI
Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg III 620 610 615 625 640 625 620 640 630 640 625 640 625 645 645 650 650 655 645 650 6670	Asp Val Pro Val IIII Ash 120 600	-~ ጥኮ~
Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp G40 625 630 645 Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 655 645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	595	d III
Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp G40 625 630 645 Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe 655 645  Asp Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys	Tyr Pro Leu Pro Gin Gin Het Arg 142 1620	-
Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg File 655 645 670 670	610 The Low Asp Gln Pro Ile Val Phe As	sp Leu
Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg File 655 645 670 670	Ser Asn Leu Ala Pro Gly Thr Leu Asp 635	640
Tan Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Led Lys	630 Fig. 7 Pho App Leu Gly A	rg Phe
Tan Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Led Lys	Leu Leu Asn Asn Leu Gly Glu Thr Pne Asp Leu Geo 6	55
Asn Cys Pro Val Asn Gly Thr Tyr Val Phe lie File 115 1670 665 670  Leu Ala Val Asn Val Pro Leu Tyr Val Asn Leu Met Lys Asn Glu Glu	645 645 Double His Met L	eu Lys
665 660 Leu Ala Val Asn Val Pro Leu Tyr Val Asn Leu Met Lys Asn Glu Glu	Nan Cys Pro Val Asn Gly Thr Tyr Val Pne 11e file 115 1670	
Leu Ala Val Asn Val Pro Leu Tyr Val Asn Leu Met Lys Ash Gra	ASII CYS 110 660 665	lu Glu
Ten var von	The Val Asp Val Pro Leu Tyr Val Asp Leu Met Lys Ash o	
	Ten var var var	

685
675  Val Leu Val Ser Ala Tyr Ala Asn Asp Gly Ala Pro Asp His Glu Thr  700  695  Cly Asp Gln Ile Trp
Val Leu Val Ser Ala Tyr Ara 1997 700 695 700 Gly Asp Gln Ile Trp
Val Leu Val Ser Ala 112 695 700 700 690 690 Ala Ser Asn His Ala Ile Leu Gln Leu Phe Gln Gly Asp Gln Ile Trp 720 715 710 715 715 710 715 715 710 715 715 710 715 715 715 715 715 715 715 715 715 715
705 Leu Arg Leu His Arg Gly Ala Ile Tyr Gly Ser Sel 119 27 735 735
725 Thr Phe Ser Gly Tyr Leu Leu Tyr Gln Asp 745 740
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Ala Leu Leu Asn Val Arg Gin 10  5  Cluster Thr Ile Val Met Thr
Ala Leu Leu Ash var Ing 10  1
The Wal Thr Thr Asn Arg Gin Thr 11e 145
Lys Thr Pro Pro Val 1122 40  35  Phe Ile Gln Thr Thr Ala Ser Thr Arg Pro Ser Val Ser Ala Pro Thr  60  55  The Rep Gln Val Gln Leu
Phe Ile Gln Thr Thi Ala 555  55  50  Ser Lys Asp Gln Val Gln Leu
Phe Ile Gin Thi The 55 55 50 Val Arg Asn Ala Met Thr Ser Ala Pro Ser Lys Asp Gln Val Gln Leu 75 70 80 Val Arg Asn Ala Met Thr Ser Ala Pro Ser Lys Asp Gln Val Gln Leu
65 Asn Asn Ser Leu Asn Giu Leu Met 272 95
Top Ile Ala Gln Pro Val Ala Thr Ala Ala 110
Pro Pro Ala Ash Tie Ala 105  100  Ser Ash Gly Thr Val Lys Lys Glu Ser Ser Ash Lys Glu Gly Ala Arg  125  120  Ser Ash Gly Thr Wet Lys
Ser Asn Gly Thr Val Lys Lys Glu Ser 125  120  125  125  127  128  129  125  120  125  125  127  128  129  129  120  120  120  120  120  120
Ser Asn Gly Thr Var 17 120  115  120  120  140  135  Met Trp Ile Asn Asp Met Lys Met Arg Ser Phe Ser Pro Thr Met Lys  140  135  135  140  135  140  135
130 Lys Glu Asp Asp Glu Pro Glu Glu Glu Glu 137 160
Val Pro Val Val Bys Clu 155  150  145  Glu Met Gly His Ala Glu Thr Tyr Ala Glu Tyr Met Pro Ile Lys Leu  175  170  165
Glu Met Gly His Ala Glu III 172 170 173 175 175 175 175 175 175 175 175 175 175
Glu Met Gly His Ald Old 170  165  Lys Ile Gly Leu Arg His Pro Asp Ala Val Val Glu Thr Ser Ser Leu  180  185  Thr Ser Ile Ser Glu
The Pro Pro Asp Val Trp Tyr Lys 1111 Set 122
Ser Ser Val IIII 110 200 200 200 195 200 Glu Thr Ile Asp Asn Gly Trp Leu Ser Ala Leu Gln Leu Glu Ala Ile 220 215 215 220 Asp Arg
Glu Thr Ile Asp Ash Gly Asp Arg
Glu Thr 11e Asp Ash 215 215 210 215 210 Thr Tyr Ala Ala Gln Gln His Glu Thr Phe Leu Pro Asn Gly Asp Arg 240 235 230 240 240 250 260 270 280 280 280 280 280 280 280 280 280 28
225 Ala Gly Asp Gly Ala Gly Val Gly Hyb 321 255
Ala Gly Phe Let 110 527 250  245  Ile Ala Gly Ile Ile Tyr Glu Asn Tyr Leu Leu Ser Arg Lys Arg Ala  260  260  260  260  260  260  260  26
11e Ala Giy 11e 120 132 265 260 260 Lvs Tyr Asp Ala Glu Arg Asp
Leu Trp Phe Ser Val Ser Asn Asp Leu Lys Tyr Asp Ala Glu Arg Asp  265  260  Leu Trp Phe Ser Val Ser Asn Asp Leu Lys Tyr Asp Ala Glu Arg Asp  285  280  280  281  285  285
Leu Trp Phe Ser var 552 280 275 280 Leu Arg Asp Ile Gly Ala Lys Asn Ile Leu Val His Ser Leu Asn Lys 300 290

Phe Lys Tyr Gly Lys Ile Ser Ser Lys His Asn Gly Ser Val Lys 320
Phe Lys Tyr Gly Lys Tie 315 310 305 305 Gly Val Ile Phe Ala Thr Tyr Ser Ser Leu Ile Gly Glu Ser Gln Ser 335 330 325
Gly Val Ile Phe Ala Thr Tyr 330  325  Gly Gly Lys Tyr Lys Thr Arg Leu Lys Gln Leu Leu His Trp Cys Gly  350  345  Gly Gly Lys Tyr Lys Thr Arg Leu Lys Gln Leu Leu His Lys Ala Lys
Gly Gly Lys Tyr Lys Thr Arg Bet 27  340  Asp Asp Phe Asp Gly Val Ile Val Phe Asp Glu Cys His Lys Ala Lys  365  360  The Lys Thr Gly Leu Ala
Asp Asp Phe Asp Gly Val Tie Val Thou 365 360 355 Asn Leu Cys Pro Val Gly Ser Ser Lys Pro Thr Lys Thr Gly Leu Ala 380 375 380
Asn Leu Cys Pro Val Gly Sel Sel 27 380 375 370 Val Leu Glu Leu Gln Asn Lys Leu Pro Lys Ala Arg Val Val Tyr Ala 395 400 Val Leu Glu Leu Gln Asn Arg
Val Leu Glu Leu Gln Asn Lys Bed 395 390 385 Ser Ala Thr Gly Ala Ser Glu Pro Arg Asn Met Ala Tyr Met Asn Arg 415 410 410 A05
Ser Ala Thr Gly Ala Ser Glu 110 1410 405 405 406 407 Leu Gly Ile Trp Gly Glu Gly Thr Pro Phe Arg Glu Phe Ser Asp Phe 430 425 430 425
Leu Gly Ile Trp Gly Glu Gly 112 425 420 420 Ile Gln Ala Val Glu Arg Arg Gly Val Gly Ala Met Glu Ile Val Ala 445 440 440 Ala Arg Gln Leu Ser Phe
Ile Gln Ala Val Glu Arg Arg 440 435  Met Asp Met Lys Leu Arg Gly Met Tyr Ile Ala Arg Gln Leu Ser Phe 460 455  Met Asp Met Lys Leu Arg Gly Met Tyr Ile Ala Arg Gln Ser Tyr
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465 An Lys Ala Val Lys Leu Trp Val 113 495
ala Ala Asp Leu Ile Asp Ala Giu Gii
Cly Cln Phe Trp Ser Ala His Gill Ala
515 The Ser Lys Val Lys Arg Val Val
530 San Gly Lys Cys Val Val Tie Gry Bot 560
545 S. T. The Leu Glu Ala Leu Glu Glu Glu Gly 527 5 575
Sor Thr Ala Lys Gly Val Leu Gin 500
The Bro Asp Arg Lys Let Tyl Co.
The Ala Pro Ser Asn Asn Ser Ser Floring Inc.
610 Lys Lys Arg Lys Gly G10 640
625 Ala Arg Lys Val Gly Gly Leu III Gry 655
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400
395 385 390 390 395 Arg Gly Arg Arg Ala Arg Gly Gln Arg Ala Gly Glu Glu Ala Gln Asp 415 405 405 410 Arg Pro Gly Asp
Arg Gly Arg Arg Pro Gly Asp
Arg Gly Arg Arg Ala May 410  405  Gly Leu Leu Pro Arg Gly Arg Asp Arg Leu Pro Leu Arg Pro Gly Asp  430  425  Arg Gly Gly His Gly
Ala Asn Gln Arg Ala Glu Arg Pro Gly Pro Pro Arg Gly Gly His Gly  445  440  Ala Asn Gln Arg Ala Glu Arg Pro Gly Pro Pro Arg His Pro
Ala Asn Gln Arg Ala Glu Arg Plo Gly 125 445
Ala Asn Gln Arg Ala 322 440 435  Pro Val Asn Ala Ser Ser Ala Pro Asp Thr Ser Pro Pro Arg His Pro 460 455  Arg Gln Phe Arg
Pro Val Asn Ala Ser Ser Mid 460
450 Ser Gln Gln Arg Gln Arg Leu 11p Arg 521 480
Arg Arg 11p val 470 470 Arg Arg Pro Pro Ala Val
Arg Arg Trp Val Ser 470 470 465 Val Gly Gly Phe Pro Pro Pro Pro Pro Ser Arg Pro Pro Ala Val 490 485
Val Gly Gly Gly The 194 490 485  Leu Leu Pro Leu Leu Arg Leu Ala Cys Ala Gly Asp Pro Gly Ala Thr 510 505 505 507 508 509 509
Leu Leu Pro Leu Leu Arg Leu Ala Cyb 122 510
Leu Leu Pro Leu leu 119 505 500 Arg Gly Glu Leu Ile  Arg Pro Gly Pro Arg Arg Pro Ala Arg Arg Pro Arg Gly Leu Arg Met
Arg Pro Gly Pro Arg Arg 520 520 520 Cly Gly Leu Arg Met
Arg Pro Gly Plo Arg 125 520 525 525 Fro Arg Arg Pro Asp Pro Ala Ala Pro Ser Glu Glu Gly Leu Arg Met 540 535 535 Fro Ala Asp Ala Ala
Fro Arg Arg Pro Asp 115 535 540 530 530 535 560 550 550 550 550 550 550 550 550 55
Glu Ser Ser Val Asp Asp Gly Ala Thr Ata The 560
Glu Ser Ser Val Asp 155 550 550 550 570 Ser Gly Glu Ala Pro Glu Ala Gly Pro Ser Pro Ser His Ser Pro Thr 575 570 570 570 570 570 570 570 570 570
Ser Gly Glu Ala Pro Glu Ala Gly 120 570
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Trp Leu Pro
595
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Glu Met Arg Lys The 10  1  1  1  1  Pro Ala Met Leu Glu Thr Leu Tyr Ser Asp Pro His Tyr Arg Ala His  25  25  20  25  20  20  20  20  20  20
Pro Ala Met Leu Giu 1112 25 25 25 25 20 Val Tyr Lys Val Leu
Pro Ala Met Hed Sta 25  20  20  25  Thr Asn Lys Asp Val Tyr Lys Val Leu  Phe Pro Asn Pro Arg Pro Asp Thr Asn Lys Asp Val Rho Glu Arg Asn  45
Phe Pro Ash Flo May 40 35 Pro Glu Ser Lys Lys Ala Pro Gly Ser Gly Ala Val Phe Glu Arg Ash 60 55 Pro Gly Leu Gln Pro
Pro Glu Ser Lys Lys Ala Pro Gly Ser Cla Pro
Fro Glu Ser Lys Lys Lys Sp 55  50  Gly Pro His Ala Ser Ser Ser Gly Val Leu Pro Leu Gly Leu Gln Pro 80  75  70  70  70  75  70  75  77  78  78  78  78  78  78  78  78
Gly Pro His Ala Ser Ser Sol 501 75
65 Star Lou Ser Lys Ser Leu Ser Ser Gin val 11p of 95
Ala pro Gly Bod 55 90 90 85 Sala Car Cys Glu Leu Ser Thr Cys
Pro Asp Pro Trp His Pro Gly Glu Gln Ser Cys Glu Leu Ser Thr Cys  110  105  105  107  108  109  109  110  100  105
100 I I I I I I I I I I I I I I I I I I
Arg Gln Gln Leu Glu Leu Ile Arg Leu Gln Met Glu Gln Met Gln Leu  125 120 120 125 120 120 125 120 120 125 120 120 120 120 120 120 120 120 120 120
115 Ala Phe Ala Pro Leu Box
Ala Met Cys His His Pro Ala Ma
Arg Gin Gin hed 525 120 120 125 125 126 Pro Leu Leu Gln Asn Gly Ala Met Cys His His Pro Ala Ala Phe Ala Pro Leu Leu 140 135 140 140 Asn Ser Asn Glu
Gln Asn Gly Ala Met Cys His His Pro Ala Ala 140 130 135 140 130 155 160
Gln Asn Gly Ala Met Cys His His Pro Ala 1140 130 130 140 Pro Thr Leu Glu Pro Ala Gln Trp Leu Ser Ile Leu Asn Ser Asn Glu 150 150

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His Leu Leu Lys Glu Lys Glu Leu Leu Ile Asp Lys Gln Arg Lys His
Ile Ser Gln Leu Glu Gln Lys Val Arg Glu Ser Glu Leu Gln Val His
Ser Ala Leu Leu Gly Arg Pro Ala Pro Phe Gly Asp Val Cys Leu Leu
Arg Leu Gln Glu Leu Gln Arg Glu Asn Thr Phe Leu Arg Ala Gln Phe
Ala Gln Lys Thr Glu Ala Leu Ser Lys Glu Lys Met Glu Leu Glu Lys
Lys Leu Ser Ala Ser Glu Val Glu Ile Gln Leu Ile Arg Glu Ser Leu
 Lys Val Thr Leu Gln Lys His Ser Glu Glu Gly Lys Lys Gln Glu Glu
 Arg Val Lys Gly Arg Asp Lys His Ile Asn Asn Leu Lys Lys Lys Cys
 Gln Lys Glu Ser Glu Gln Asn Arg Glu Lys Gln Gln Arg Ile Glu Thr
 Leu Glu Arg Tyr Leu Ala Asp Leu Pro Thr Leu Glu Asp His Gln Lys
 Gln Thr Glu Gln Leu Lys Asp Ala Glu Leu Lys Asn Thr Glu Leu Gln
  Glu Arg Val Ala Glu Leu Glu Thr Leu Leu Glu Asp Thr Gln Ala Thr
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          355
  Asp Leu Ser Ser Ala Arg His Arg
      370
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         <211> 160
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   Thr Thr Ala Asn Arg Thr Gln Ser Leu Asn Tyr Gly Cys Ile Val Glu
   Asn Pro Gln Thr His Glu Val Leu His Tyr Val Glu Lys Pro Ser Thr
    Phe Ile Ser Asp Ile Ile Asn Cys Gly Ile Tyr Leu Phe Ser Pro Glu
    Ala Leu Lys Pro Leu Arg Asp Val Phe Gln Arg Asn Gln Gln Asp Gly
    Gln Leu Glu Asp Ser Pro Gly Leu Trp Pro Gly Ala Gly Thr Ile Arg
    Leu Glu Gln Asp Val Phe Ser Ala Leu Ala Gly Gln Gly Gln Ile Tyr
    Val His Leu Thr Asp Gly Ile Trp Ser Gln Ile Lys Ser Ala Gly Ser
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160 155 150 145 <210> 190 <211> 146 <212> PRT <213> Homo sapien Met Asp Pro Arg Ala Ser Leu Leu Leu Leu Gly Asn Val Tyr Ile His Pro Thr Ala Lys Val Ala Pro Ser Ala Val Leu Gly Pro Asn Val Ser Ile Gly Lys Gly Val Thr Val Gly Glu Gly Val Arg Leu Arg Glu Ser Ile Val Leu His Gly Ala Thr Leu Gln Glu His Thr Cys Val Leu His Ser Ile Val Gly Trp Gly Ser Thr Val Gly Arg Trp Ala Arg Val Glu Gly Thr Pro Ser Asp Pro Asn Pro Asn Asp Pro Arg Ala Arg Met Asp Ser Glu Ser Leu Phe Lys Asp Gly Lys Leu Leu Pro Ala Ile Thr Ile Leu Gly Cys Arg Val Arg Ile Pro Ala Glu Val Leu Ile Leu Asn Ser Ile Val Leu Pro His Lys Glu Leu Ser Arg Ser Phe Thr Asn Gln Ile 130 Ile Leu 145 <210> 191 <211> 704 <212> PRT <213> Homo sapien Glu Gly Gly Cys Ala Ala Gly Arg Gly Arg Glu Leu Glu Pro Glu Leu Glu Pro Gly Pro Gly Pro Gly Ser Ala Leu Glu Pro Gly Glu Glu Phe Glu Ile Val Asp Arg Ser Gln Leu Pro Gly Pro Gly Asp Leu Arg Ser Ala Thr Arg Pro Arg Ala Ala Glu Gly Trp Ser Ala Pro Ile Leu Thr Leu Ala Arg Arg Ala Thr Gly Asn Leu Ser Ala Ser Cys Gly Ser Ala Leu Arg Ala Ala Gly Leu Gly Gly Gly Asp Ser Gly Asp Gly Thr Ala Arg Ala Ala Ser Lys Cys Gln Met Met Glu Glu Arg Ala Asn Leu Met His Met Met Lys Leu Ser Ile Lys Val Leu Leu Gln Ser Ala Leu Ser Leu Gly Arg Ser Leu Asp Ala Asp His Ala Pro Leu Gln Gln Phe Phe Val Val Met Glu His Cys Leu Lys His Gly Leu Lys Val Lys Lys

150	155	160
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Ser Phe Ile Gly GIn ASI	170	val Arg Asn
Glu Lys Leu Cys Pro Glu	Ala Ser Asp Ile Ala Thr Ser	190
Leu Pro Glu Leu Lys Thr	185 Ala Val Gly Arg Gly Arg Ala 200	o val Leu Ile
Leu Ala Leu Met Gln Lys	Lys Leu Ala Asp Tyr Leu Lys 220 215	n Ala Leu Met
210	215 Ser Glu Phe Tyr Glu Pro Gl 235	240
23 225	Val Ile Val Gly Leu Leu Va 250	l Gly Leu Asn
Met Glu Glu Glu Gly Me 245	250	eu Asp Ser Gln
Val Leu Asp Ala Asn Le	250 1 Cys Leu Lys Gly Glu Asp Le 265	270
260	265 Ser Leu Tyr Leu Lys Asp Va 280 280	al Gin Asp Leu
Val Gly Val Tie ASP 11	280 Zen Asp Val Le	eu Asp Gln Lys
Asp Gly Gly Lys Glu H	280 s Glu Arg Ile Thr Asp Val Le 300 295	
290	295 u Asn Arg His Leu Ser Cys T 0 315	hr Val Gly Asp 320
Asn Tyr Val Glu Glu B	0 315	or Lys Leu Gln
305	0 p Gly Leu Glu Lys Thr Asn S 330	335
325	330 The Arm Arm The Cys Ser I	eu Gln Glu Glu
Glu Glu Leu Ser Ala A	a Thr Asp Arg Ile Cys Ser I	350
340	345 Lu Gln Asn Glu Leu Ile Arg G 360	Hu Arg Ser Gra
Gln Gln Gln Hed 1119	360	Glu Leu Glu Thr
Lys Ser Val Glu Ile S	360 hr Lys Gln Asp Thr Lys Val ( 380	
370	375 ln Gly Leu Asp Glu Met Tyr	Ser Asp Val 11P
Tyr Lys Gin IIII Arg	90 395 Hu Lys Lys Val Arg Leu Glu 410	Leu Glu Lys Glu
Lys Gln Leu Lys Glu	lu Lys Lys var Mag = 410	415
405	31y Met Lys Thr Glu Met Glu 425	Ile Ala Met 175
Leu Giu Leu Gin 220	425	Leu Val Ala Leu
Leu Leu Glu Lys Asp	425 Thr His Glu Lys Gln Asp Thr 440	445
435	440 Glu Val Lys Ala Ile Asn Leu 455	Gln Met Pne HIS
Arg Gln Gln Led Glu	455 Gla Cla Lys	Asn Glu Ala Ile
Lys Ala Gln Asn Ala	455 Glu Ser Ser Leu Gln Gln Lys 475	480
465	470 Lys Thr Asn Gln Val Met Ser 490	Ser Met Lys Gin
Thr Ser Phe Glu Gly	490	. Cln Glv Ala Glu
Met Glu Glu Arg Lev	Gln His Ser Glu Arg Ala Arg	510
500	505 Leu Gln Gln Glu Leu Gly Gly 520	Arg Ile Gly Ala
Glu Arg Ser His Lys	520 Cln Cur	525 Ser Ser Leu Glü
Jon Gln Leu Gln Lei	520 Ser Gln Leu His Glu Gln Cys 535	)
530	535	a Leu Gln Arg Glu
Lys Glu Leu Lys Se	535 Glu Lys Glu Gln Arg Gln Al	56
545	Asp Thr Ser Ser Leu Leu Ar	575
Leu Gin His Gid by	570 Lys Lys Glu Leu Arg Glu Le	u Gln Asp Glu Ly
	Tare Tite Clin Leu Ary Gra no	

590
580  580  580  Ala Glu Leu Gln Lys Ile Cys Glu Glu Gln Glu Gln Ala Leu Gln Glu 605  600  600  601  605  605
Ala Glu Leu Gin Lys 600  595  Met Gly Leu His Leu Ser Gln Ser Lys Leu Lys Met Glu Asp Ile Lys  620  615  620  615
Met Gly Leu His Leu Sel Gli 505 620 620 615 610 Fig. Nia Trp Leu Lys Asp Asp Glu
Met Gly Leu HIS hed 615 610 610 610 Glu Val Asn Gln Ala Leu Lys Gly His Ala Trp Leu Lys Asp Asp Glu 630 630 630 630 630 630 630 630 630 630
625 Glu Lys Glu Phe Sel 11c 555
Gys Arg Asn Cys Gly His Ile Phe Cys Ash Inc 670
Lys His His Cys Arg Val 665  660  Ser Asn Glu Leu Ala Leu Pro Ser Tyr Pro Lys Pro Val Arg Val Cys  685  680  Ger Ser Thr Ala Ser
Ser Asn Glu Leu Ala Leu 110 680 685 680 675 680 Ser Ser Thr Ala Ser
Ser Asn Glu Leu Ard 200 680 685  675 680 Asp Ser Cys His Thr Leu Leu Leu Gln Arg Cys Ser Ser Thr Ala Ser  690 695
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Arg Ala Gly Ala Sol 10  1  1  1  1  10  1  1  11  10  11  11  10
are Cla Leu Lys Val Leu Ser His Asn Leu 191 1111 145
Ser Lys Ser Gin Hod 27  40  35  His Ile Pro His Asp Pro Val Ala Leu Glu Glu His Phe Arg Asp Asp  60  55  60  75  75  75  75  75  75  75  75  75  7
His Ile Pro his Asp 55 55 50 Asp Asp Gly Pro Val Ser Ser Gln Gly Tyr Met Pro Tyr Leu Asn Lys 80 75 75 75 75 76 77 78 78 78 78 78 78 78 78 78 78 78 78
Asp Asp Gly Pro Val Sel Sel Strain 75 75 70 70 70 Phe Val Lys Glu His Phe
Asp Asp Gly Flo Val 70  70  65  Tyr Ile Leu Asp Lys Val Glu Glu Glu Gly Ala Phe Val Lys Glu His Phe 95  78  79  70  70  70  70  70  70  70  70  70
The Cyc Trp Thr Leu Thr Ala Lys Lys Ash 191 Alg 110
Age Age Ser Met Leu Ser Asn Gln Asp Ala Phe Alg 200 125
Ser Asn Gly Ash Ser 120  115  Cys Leu Phe Asn Phe Leu Ser Glu Asp Lys Tyr Pro Leu Ile Met Val  135  140  135  140  135  140  140  135
Cys Leu Phe Ash File Hou 135 140  130 135 150 150  Pro Asp Glu Val Glu Tyr Leu Leu Lys Lys Val Leu Ser Ser Met Ser 160  150 150 155 160 Ala Gln Glu Ala
Pro Asp Glu Val Glu Tyr Leu Leu Lys 155 150 150 155 Ala Gln Glu Ala
Pro Asp Glu Val Glu 175  150  145  Leu Glu Val Ser Leu Gly Glu Leu Glu Glu Leu Leu Ala Gln Glu Ala  175  170  170  185  185  185  186  186  187  188  188  188  188  188
The Cla Thr Thr Gly Gly Leu Ser Val Trp Gin 190
Gln Val Ala Gln Thr Leu  180  180  Leu Phe Asn Ser Gly Arg Cys Leu Arg Gly Val Gly Arg Asp Thr Leu  205  200  The Cln Asp Val Leu
Leu Phe Asn Ser Gly Ala 200  195  Ser Met Ala Ile His Glu Val Tyr Gln Glu Leu Ile Gln Asp Val Leu  220  215  220  215  Ang Arg Asn Trp Ala
Ser Met Ala Ile His Giu vai 191 220 215 210 210 210 210 210 210 210 210 210 210
Ser Met Ala Tie His 3215 215 210 215 Lys Gln Gly Tyr Leu Trp Lys Arg Gly His Leu Arg Arg Asn Trp Ala 240 235 230 230 237 240 257 258 258 259 259 250 250 250 250 250 250 250 250 250 250
Lys Gin Giy Tyr Edd 230 230 225 230 Clu Arg Trp Phe Gin Leu Gin Pro Ser Cys Leu Cys Tyr Phe Gly Ser 255 245
230

Glu Glu Cys Lys Glu Lys Arg Gly Ile Ile Pro Leu Asp Ala His Cys Cys Val Glu Val Leu Pro Asp Arg Asp Gly Lys Arg Cys Met Phe Cys Val Lys Thr Ala Thr Arg Thr Tyr Glu Met Ser Ala Ser Asp Thr Arg Gln Arg Gln Glu Trp Thr Ala Ala Ile Gln Met Ala Ile Arg Leu Gln 310 Ala Glu Gly Lys Thr Ser Leu His Lys Asp Leu 325 <210> 193 <211> 475 <212> PRT <213> Homo sapien Lys Asn Ser Pro Leu Leu Ser Val Ser Ser Gln Thr Ile Thr Lys Glu Asn Asn Arg Asn Val His Leu Glu His Ser Glu Gln Asn Pro Gly Ser Ser Ala Gly Asp Thr Ser Ala Ala His Gln Val Val Leu Gly Glu Asn Leu Ile Ala Thr Ala Leu Cys Leu Ser Gly Ser Gln Ser Asp Leu Lys Asp Val Ala Ser Thr Ala Gly Glu Glu Gly Asp Thr Ser Leu Arg Glu Ser Leu His Pro Val Thr Arg Ser Leu Lys Ala Gly Cys His Thr Lys Gln Leu Ala Ser Arg Asn Cys Ser Glu Glu Lys Ser Pro Gln Thr Ser Ile Leu Lys Glu Gly Asn Arg Asp Thr Ser Leu Asp Phe Arg Pro Val Val Ser Pro Ala Asn Gly Val Glu Gly Val Arg Val Asp Gln Asp Asp Asp Gln Asp Ser Ser Ser Leu Lys Leu Ser Gln Asn Ile Ala Val Gln Thr Asp Phe Lys Thr Ala Asp Ser Glu Val Asn Thr Asp Gln Asp Ile Glu Lys Asn Leu Asp Lys Met Met Thr Glu Arg Thr Leu Leu Lys Glu Arg Tyr Gln Glu Val Leu Asp Lys Gln Arg Gln Val Glu Asn Gln Leu Gln Val Gln Leu Lys Gln Leu Gln Gln Arg Arg Glu Glu Glu Met Lys Asn His Gln Glu Ile Leu Lys Ala Ile Gln Asp Val Thr Ile Lys Arg Glu Glu Thr Lys Lys Ile Glu Lys Glu Lys Glu Phe Leu Gln Lys Glu Gln Asp Leu Lys Ala Glu Ile Glu Lys Leu Cys Glu Lys Gly Arg Arg Glu Val Trp Glu Met Glu Leu Asp Arg Leu Lys Asn Gln Asp Gly Glu Ile Asn Arg Asn Ile Met Glu Glu Thr Glu Arg Ala

290 295 300  Trp Lys Ala Glu Ile Leu Ser Leu Glu Ser Arg Lys Glu Leu Leu Val 320 305  Leu Lys Leu Glu Glu Ala Glu Lys Glu Ala Glu Leu His Leu Thr Tyr 335  Leu Lys Ser Thr Pro Pro Thr Leu Glu Thr Val Arg Ser Lys Gln Glu 340  Trp Glu Thr Arg Leu Asn Gly Val Arg Ile Met Lys Lys Asn Val Arg 355  Asp Gln Phe Asn Ser His Ile Gln Leu Val Arg Asn Gly Ala Lys Leu 370  Ser Ser Leu Pro Gln Ile Pro Thr Pro Thr Leu Pro Pro Pro Ser 380 380 390  Glu Thr Asp Phe Met Leu Gln Val Phe Gln Pro Ser Pro Ser Leu Ala 405  Pro Arg Met Pro Phe Ser Ile Gly Gln Val Thr Met Pro Met Val Met 420  Pro Ser Ala Asp Pro Arg Ser Leu Ser Phe Pro Ile Leu Asn Pro Ala
Pro Ser Ala Asp Pro Arg 304 440 445  Leu Ser Gln Pro Ser Gln Pro Ser Ser Pro Leu Pro Gly Ser His Gly  455 460  Arg Asn Ser Pro Gly Leu Gly Ser Leu Val Ser  475  465
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Gln Lys Met Ile Leu Glu Leu Phe Ser Lys Val 125 120 115 Ser Phe Ile Arg Ser Gln Asn Lys Glu Asp Tyr Ala Gly Leu Lys Glu 135 130 130 140 160
Glu Phe Arg Lys Glu File 112 1 155  145  Lys Thr Thr Phe Phe Gly Gly Asn Ser Ile Ser Met Ile Asp Tyr Leu 175  165  Ile Trp Pro Trp Phe Glu Arg Leu Glu Ala Met Lys Leu Asn Glu Cys 180

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Val Asp His Thr Pro Lys Leu Lys Leu Trp Met Ala Ala Met Lys Glu
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  Arg Ala Glu Glu Gly Lys Gly Pro Ser Lys Ala Gln Arg Gly Ser Leu
  Glu His Met Lys Leu Ile Leu Arg Asp Lys Glu Lys Glu Val Glu Cys
  Gln Gln Glu His Ile His Glu Leu Gln Glu Leu Lys Asp Gln Leu Glu
  Gln Gln Leu Gln Gly Leu His Arg Lys Val Gly Glu Thr Ser Leu Leu
  Leu Ser Gln Arg Glu Gln Glu Ile Val Val Leu Gln Gln Gln Leu Gln
  Glu Ala Arg Glu Gln Gly Glu Leu Lys Glu Gln Ser Leu Gln Ser Gln
   Leu Asp Glu Ala Gln Arg Ala Leu Ala Gln
                           135
       130
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    Asp Met Leu Thr Glu Leu Ala Asn Phe Glu Lys Asn Val Ser Gln Ala
    Ile His Lys Tyr Asn Ala Tyr Arg Lys Ala Ala Ser Val Ile Ala Lys
    Tyr Pro His Lys Ile Lys Ser Gly Ala Glu Ala Lys Lys Leu Pro Gly
     Val Gly Thr Lys Ile Ala Glu Lys Ile Asp Glu Phe Leu Ala Thr Gly
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     Ile Asn Phe Leu Thr Arg
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Tyr Ser Glu Cys Glu Thr Phe Thr Asp Glu Asp Thr Ser Thr Leu Val
His Pro Glu Leu Gln Pro Glu Gly Asp Ala Asp Ser Ala Gly Gly Ser
Ala Val Pro Ser Glu Cys Leu Asp Ala Met Glu Glu Pro Asp His Gly
Ala Leu Leu Leu Pro Gly Arg Pro His Pro His Gly Gln Ser Val
 Ile Thr Val Ile Gly Gly Glu Glu His Phe Glu Asp Tyr Gly Glu Gly
 Ser Glu Ala Glu Leu Ser Pro Glu Thr Leu Cys Asn Gly Gln Leu Gly
 Cys Ser Asp Pro Ala Phe Leu Thr Pro Ser Pro Thr Lys Arg Leu Ser
 Ser Lys Lys Val Ala Arg Tyr Leu His Gln
                          135
        <210> 198
        <211> 100
        <212> PRT
        <213> Homo sapien
  Met Gly Asp Val Lys Asn Phe Leu Tyr Ala Trp Cys Gly Lys Arg Lys
  Met Thr Pro Ser Tyr Glu Ile Arg Ala Val Gly Asn Lys Asn Arg Gln
   Lys Phe Met Cys Glu Val Gln Val Glu Gly Tyr Asn Tyr Thr Gly Met
   Gly Asn Ser Thr Asn Lys Lys Asp Ala Gln Ser Asn Ala Ala Arg Asp
   Phe Val Asn Tyr Leu Val Arg Ile Asn Glu Ile Lys Ser Glu Glu Val
   Pro Ala Phe Gly Val Ala Ser Pro Pro Pro Leu Thr Asp Thr Pro Asp
                    85
    Thr Thr Ala Asn
                100
          <210> 199
          <211> 127
          <212> PRT
          <213> Homo sapien
    Met Val Lys Glu Thr Thr Tyr Tyr Asp Val Leu Gly Val Lys Pro Asn
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Ala Thr Gln Glu Glu Leu Lys Lys Ala Tyr Arg Lys Leu Ala Leu Lys
Tyr His Pro Asp Lys Asn Pro Asn Glu Gly Glu Lys Phe Lys Gln Ile
Ser Gln Ala Tyr Glu Val Leu Ser Asp Ala Lys Lys Arg Glu Leu Tyr
Asp Lys Gly Gly Glu Gln Ala Ile Lys Glu Gly Gly Ala Gly Gly
Phe Gly Ser Pro Met Asp Ile Phe Asp Met Phe Phe Gly Gly Gly
 Arg Met Gln Arg Glu Arg Arg Gly Lys Asn Val Val His Gln Leu Ser
 Val Thr Leu Glu Asp Leu Tyr Asn Gly Ala Thr Arg Lys Leu Ala
       <210> 200
       <211> 90
       <212> PRT
       <213> Homo sapien
  Met Ala Cys Pro Leu Asp Gln Ala Ile Gly Leu Leu Val Ala Ile Phe
  His Lys Tyr Ser Gly Arg Glu Gly Asp Lys His Thr Leu Ser Lys Lys
  Glu Leu Lys Glu Leu Ile Gln Lys Glu Leu Thr Ile Gly Ser Lys Leu
  Gln Asp Ala Glu Ile Ala Arg Leu Met Glu Asp Leu Asp Arg Asn Lys
  Asp Gln Glu Val Asn Phe Gln Glu Tyr Val Thr Phe Leu Gly Ala Leu
                       70
   Ala Leu Ile Tyr Asn Glu Ala Leu Lys Gly
                   85
         <210> 201
         <211> 120
          <212> PRT
          <213> Homo sapien
    Met Glu Thr Pro Ser Gln Arg Arg Ala Thr Arg Ser Gly Ala Gln Ala
    Ser Ser Thr Pro Leu Ser Pro Thr Arg Ile Thr Arg Leu Gln Glu Lys
    Glu Asp Leu Gln Glu Leu Asn Asp Arg Leu Ala Val Tyr Ile Asp Arg
    Val Arg Ser Leu Glu Thr Glu Asn Ala Gly Leu Arg Leu Arg Ile Thr
     Glu Ser Glu Glu Val Val Ser Arg Glu Val Ser Gly Ile Lys Ala Ala
     Tyr Glu Ala Glu Leu Gly Asp Ala Arg Lys Thr Leu Asp Ser Val Ala
     Lys Glu Arg Ala Arg Leu Gln Leu Glu Leu Ser Lys Val Arg Glu Glu
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Phe Lys Glu Leu Lys Ala Arg Asn
              115
            <210> 202
             <211> 177
             <212> PRT
             <213> Homo sapien
       Met Ala Ala Gly Val Glu Ala Ala Ala Glu Val Ala Ala Thr Glu Ile
       Lys Met Glu Glu Glu Ser Gly Ala Pro Gly Val Pro Ser Gly Asn Gly
       Ala Pro Gly Pro Lys Gly Glu Gly Glu Arg Pro Ala Gln Asn Glu Lys
        Arg Lys Glu Lys Asn Ile Lys Arg Gly Gly Asn Arg Phe Glu Pro Tyr
        Ala Asn Pro Thr Lys Arg Tyr Arg Ala Phe Ile Thr Asn Ile Pro Phe
Asp Val Lys Trp Gln Ser Leu Lys Asp Leu Val Lys Glu Lys Val Gly
        Glu Val Thr Tyr Val Glu Leu Leu Met Asp Ala Glu Gly Lys Ser Arg
         Gly Cys Ala Val Val Glu Phe Lys Met Glu Glu Ser Met Lys Lys Ala
ļå
         Ala Glu Val Leu Asn Lys His Ser Leu Ser Gly Arg Pro Leu Lys Val
IJ
IJ
         Lys Glu Asp Pro Asp Gly Glu His Ala Arg Arg Ala Met Gln Lys Ala
31
 Gly Arg Leu Gly Ser Thr Val Phe Val Ala Asn Leu Asp Tyr Lys Val
 M
 1.4
                          165
 14
          Gly
                <210> 203
                <211> 164
                <212> PRT
                <213> Homo sapien
          Met Arg Leu Ala Val Gly Ala Leu Leu Val Cys Ala Val Leu Gly Leu
           Cys Leu Ala Val Pro Asp Lys Thr Val Arg Trp Cys Ala Val Ser Glu
           His Glu Ala Thr Lys Cys Gln Ser Phe Arg Asp His Met Lys Ser Val
           Ile Pro Ser Asp Gly Pro Ser Val Ala Cys Val Lys Lys Ala Ser Tyr
           Leu Asp Cys Ile Arg Ala Ile Ala Ala Asn Glu Ala Asp Ala Val Thr
            Leu Asp Ala Gly Leu Val Tyr Asp Ala Tyr Leu Ala Pro Asn Asn Leu
            Lys Pro Val Val Ala Glu Phe Tyr Gly Ser Lys Glu Asp Pro Gln Thr
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Phe Tyr Tyr Ala Val Ala Val Lys Lys Asp Ser Gly Phe Gln Met
Asn Gln Leu Arg Gly Lys Lys Ser Cys His Thr Gly Leu Gly Arg Ser
Ala Gly Trp Asn Ile Pro Ile Gly Leu Leu Tyr Cys Asp Leu Pro Glu
145
Pro Arg Lys Pro
       <210> 204
       <211> 241
       <212> PRT
       <213> Homo sapien
 Met Ser Gly Glu Ser Ala Arg Ser Leu Gly Lys Gly Ser Ala Pro Pro
  Gly Pro Val Pro Glu Gly Ser Ile Arg Ile Tyr Ser Met Arg Phe Cys
  Pro Phe Ala Glu Arg Thr Arg Leu Val Leu Lys Ala Lys Gly Ile Arg
  His Glu Val Ile Asn Ile Asn Leu Lys Asn Lys Pro Glu Trp Phe Phe
  Lys Lys Asn Pro Phe Gly Leu Val Pro Val Leu Glu Asn Ser Gln Gly
   Gln Leu Ile Tyr Glu Ser Ala Ile Thr Cys Glu Tyr Leu Asp Glu Ala
   Tyr Pro Gly Lys Lys Leu Leu Pro Asp Asp Pro Tyr Glu Lys Ala Cys
   Gln Lys Met Ile Leu Glu Leu Phe Ser Lys Val Pro Ser Leu Val Gly
   Ser Phe Ile Arg Ser Gln Asn Lys Glu Asp Tyr Asp Gly Leu Lys Glu
   Glu Phe Arg Lys Glu Phe Thr Lys Leu Glu Glu Val Leu Thr Asn Lys
    Lys Thr Thr Phe Phe Gly Gly Asn Ser Ile Ser Met Ile Asp Tyr Leu
    Ile Trp Pro Trp Phe Glu Arg Leu Glu Ala Met Lys Leu Asn Glu Cys
    Val Asp His Thr Pro Lys Leu Lys Leu Trp Met Ala Ala Met Lys Glu
     Asp Pro Thr Val Ser Ala Leu Leu Thr Ser Glu Lys Asp Trp Gln Gly
     Phe Leu Glu Leu Tyr Leu Gln Asn Ser Pro Glu Ala Cys Asp Tyr Gly
     Leu
            <210> 205
```

<211> 160

<212> PRT

<213> Homo sapien

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Met Gln Ile Phe Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu
Val Glu Pro Ser Asp Thr Ile Glu Asn Val Lys Ala Lys Ile Gln Asp
Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys
Gln Leu Glu Asp Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu
 Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly Met Gln Ile Phe
 Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu Val Glu Pro Ser
 Asp Thr Ile Glu Asn Val Lys Ala Lys Ile Gln Asp Lys Glu Gly Ile
 Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys Gln Leu Glu Asp
  Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu Ser Thr Leu His
  Leu Val Leu Arg Leu Arg Gly Gly Met Gln Ile Phe Val Lys Thr Leu
  145
        <210> 206
         <211> 197
         <212> PRT
         <213> Homo sapien
   Thr Ser Pro Ser Glu Ala Cys Ala Pro Leu Leu Ile Ser Leu Ser Thr
   Leu Ile Tyr Asn Gly Ala Leu Pro Cys Gln Cys Asn Pro Gln Gly Ser
   Leu Ser Ser Glu Cys Asn Pro His Gly Gly Gln Cys Leu Cys Lys Pro
    Gly Val Val Gly Arg Arg Cys Asp Leu Cys Ala Pro Gly Tyr Tyr Gly
    Phe Gly Pro Thr Gly Cys Gln Gly Ala Cys Leu Gly Cys Arg Asp His
    Thr Gly Gly Glu His Cys Glu Arg Cys Ile Ala Gly Phe His Gly Asp
    Pro Arg Leu Pro Tyr Gly Gly Gln Cys Arg Pro Cys Pro Cys Pro Glu
     Gly Pro Gly Ser Gln Arg His Phe Ala Thr Ser Cys His Gln Asp Glu
     Tyr Ser Gln Gln Ile Val Cys His Cys Arg Ala Gly Tyr Thr Gly Leu
     Arg Cys Glu Ala Cys Ala Pro Gly His Phe Gly Asp Pro Ser Arg Pro
     Gly Gly Arg Cys Gln Leu Cys Glu Cys Ser Gly Asn Ile Asp Pro Met
     Asp Pro Asp Ala Cys Asp Pro His Thr Gly Gln Cys Leu Arg Cys Leu
                  180
      His His Thr Glu Gly
              195
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<210> 207
     <211> 175
     <212> PRT
     <213> Homo sapien
Ile Ile Arg Gln Gln Gly Leu Ala Ser Tyr Asp Tyr Val Arg Arg
Leu Thr Ala Glu Asp Leu Phe Glu Ala Arg Ile Ile Ser Leu Glu Thr
Tyr Asn Leu Leu Arg Glu Gly Thr Arg Ser Leu Arg Glu Ala Leu Glu
Ala Glu Ser Ala Trp Cys Tyr Leu Tyr Gly Thr Gly Ser Val Ala Gly
Val Tyr Leu Pro Gly Ser Arg Gln Thr Leu Ser Ile Tyr Gln Ala Leu
 Lys Lys Gly Leu Leu Ser Ala Glu Val Ala Arg Leu Leu Glu Ala
 Gln Ala Ala Thr Gly Phe Leu Leu Asp Pro Val Lys Gly Glu Arg Leu
 Thr Val Asp Glu Ala Val Arg Lys Gly Leu Val Gly Pro Glu Leu His
 Asp Arg Leu Leu Ser Ala Glu Arg Ala Val Thr Gly Tyr Arg Asp Pro
 Tyr Thr Glu Gln Thr Ile Ser Leu Phe Gln Ala Met Lys Lys Glu Leu
  Ile Pro Thr Glu Glu Ala Leu Arg Leu Trp Met Pro Ser Trp Pro
                  165
        <210> 208
        <211> 177
        <212> PRT
        <213> Homo sapien
   Met Ala Ala Gly Val Glu Ala Ala Glu Val Ala Ala Thr Glu Ile
   Lys Met Glu Glu Ser Gly Ala Pro Gly Val Pro Ser Gly Asn Gly
   Ala Pro Gly Pro Lys Gly Glu Gly Glu Arg Pro Ala Gln Asn Glu Lys
   Arg Lys Glu Lys Asn Ile Lys Arg Gly Gly Asn Arg Phe Glu Pro Tyr
    Ala Asn Pro Thr Lys Arg Tyr Arg Ala Phe Ile Thr Asn Ile Pro Phe
    Asp Val Lys Trp Gln Ser Leu Lys Asp Leu Val Lys Glu Lys Val Gly
    Glu Val Thr Tyr Val Glu Leu Leu Met Asp Ala Glu Gly Lys Ser Arg
    Gly Cys Ala Val Val Glu Phe Lys Met Glu Glu Ser Met Lys Lys Ala
    Ala Glu Val Leu Asn Lys His Ser Leu Ser Gly Arg Pro Leu Lys Val
                             135
         130
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Lys Glu Asp Pro Asp Gly Glu His Ala Arg Arg Ala Met Gln Lys Val

Met Ala Thr Thr Gly Gly Met Gly Met Gly Pro Gly Pro Gly Met

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Ile
             <210> 209
             <211> 196
             <212> PRT
             <213> Homo sapien
       Asp Leu Gln Asp Met Phe Ile Val His Thr Ile Glu Glu Ile Glu Gly
        Leu Ile Ser Ala His Asp Gln Phe Lys Ser Thr Leu Pro Asp Ala Asp
        Arg Glu Arg Glu Ala Ile Leu Ala Ile His Lys Glu Ala Gln Arg Ile
        Ala Glu Ser Asn His Ile Lys Leu Ser Gly Ser Asn Pro Tyr Thr Thr
Val Thr Pro Gln Ile Ile Asn Ser Lys Trp Glu Lys Val Gln Gln Leu
         Val Pro Lys Arg Asp His Ala Leu Leu Glu Glu Gln Ser Lys Gln Gln
         Ser Asn Glu His Leu Arg Arg Gln Phe Ala Ser Gln Ala Asn Val Val
Gly Pro Trp Ile Gln Thr Lys Met Glu Glu Ile Gly Arg Ile Ser Ile
W
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         Glu Met Asn Gly Thr Leu Glu Asp Gln Leu Ser His Leu Lys Gln Tyr
 M
          Glu Arg Ser Ile Val Asp Tyr Lys Pro Asn Leu Asp Leu Leu Glu Gln
 1,2
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          Gln His Gln Leu Ile Gln Glu Ala Leu Ile Phe Asp Asn Lys His Thr
          Asn Tyr Thr Met Glu His Ile Arg Val Gly Trp Glu Gln Leu Leu Thr
                      180
           Thr Ile Ala Arg
                   195
                 <210> 210
                 <211> 156
                 <212> PRT
                 <213> Homo sapien
           Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val Ala Glu Gly Lys Glu
            Val Leu Leu Ala His Asn Leu Pro Gln Asn Arg Ile Gly Tyr Ser
            Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Ser Leu Ile Val Gly Tyr
            Val Ile Gly Thr Gln Gln Ala Thr Pro Gly Pro Ala Tyr Ser Gly Arg
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Glu Thr Ile Tyr Pro Asn Ala Ser Leu Leu Ile Gln Asn Val Thr Gln
Asn Asp Thr Gly Phe Tyr Thr Leu Gln Val Ile Lys Ser Asp Leu Val
Asn Glu Glu Ala Thr Gly Gln Phe His Val Tyr Pro Glu Leu Pro Lys
Pro Ser Ile Ser Ser Asn Asn Ser Asn Pro Val Glu Asp Lys Asp Ala
Val Ala Phe Thr Cys Glu Pro Glu Val Gln Asn Thr Thr Tyr Leu Trp
                       135
 Trp Val Asn Gly Gln Ser Leu Pro Val Ser Pro Lys
       <210> 211
       <211> 92
       <212> PRT
       <213> Homo sapien
  Met Glu Ser Pro Ser Ala Pro Pro His Arg Trp Cys Ile Pro Trp Gln
  Arg Leu Leu Thr Ala Ser Leu Leu Thr Phe Trp Asn Pro Pro Thr
  Thr Ala Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val Ala Glu Gly
  Lys Glu Val Leu Leu Val His Asn Leu Pro Gln His Leu Phe Gly
  Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Arg Gln Ile Ile
   Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr Pro Gly
                      70
                   85
         <210> 212
         <211> 142
         <212> PRT
         <213> Homo sapien
    Glu Lys Gln Lys Asn Lys Glu Phe Ser Gln Thr Leu Glu Asn Glu Lys
    Asn Thr Leu Leu Ser Gln Ile Ser Thr Lys Asp Gly Glu Leu Lys Met
    Leu Gln Glu Val Thr Lys Met Asn Leu Leu Asn Gln Gln Ile Gln
    Glu Glu Leu Ser Arg Val Thr Lys Leu Lys Glu Thr Ala Glu Glu Glu
    Lys Asp Asp Leu Glu Glu Arg Leu Met Asn Gln Leu Ala Glu Leu Asn
     Gly Ser Ile Gly Asn Tyr Cys Gln Asp Val Thr Asp Ala Gln Ile Lys
     Asn Glu Leu Leu Glu Ser Glu Met Lys Asn Leu Lys Lys Cys Val Ser
     Glu Leu Glu Glu Lys Gln Gln Leu Val Lys Glu Lys Thr Lys Val
             115
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Glu Ser Glu Ile Arg Lys Glu Tyr Leu Glu Lys Ile Gln Gly
      <210> 213
      <211> 142
      <212> PRT
      <213> Homo sapien
 Gly Gly Tyr Gly Gly Tyr Gly Gly Val Leu Thr Ala Ser Asp Gly
 Leu Leu Ala Gly Asn Glu Lys Leu Thr Met Gln Asn Leu Asn Asp Arg
 Leu Ala Ser Tyr Leu Asp Lys Val Arg Ala Leu Glu Ala Ala Asn Gly
 Glu Leu Glu Val Lys Ile Arg Asp Trp Tyr Gln Lys Gln Gly Pro Gly
  Pro Ser Arg Asp Tyr Ser His Tyr Tyr Thr Thr Ile Gln Asp Leu Arg
  Asp Lys Ile Leu Gly Ala Thr Ile Glu Asn Ser Arg Ile Val Leu Gln
  Ile Asp Asn Ala Arg Leu Ala Ala Asp Asp Phe Arg Thr Lys Phe Glu
  Thr Glu Gln Ala Leu Arg Met Ser Val Glu Ala Asp Ile Asn Gly Leu
  Arg Arg Val Leu Asp Glu Leu Thr Leu Ala Arg Thr Asp Leu
       130
         <210> 214
         <211> 129
         <212> PRT
         <213> Homo sapien
    Val Met Arg Val Asp Phe Asn Val Pro Met Lys Asn Asn Gln Ile Thr
    Asn Asn Gln Arg Ile Lys Ala Ala Val Pro Ser Ile Lys Phe Cys Leu
    Asp Asn Gly Ala Lys Ser Val Val Leu Met Ser His Leu Gly Arg Pro
    Asp Gly Val Pro Met Pro Asp Lys Tyr Ser Leu Glu Pro Val Ala Val
    Glu Leu Arg Ser Leu Leu Gly Lys Asp Val Leu Phe Leu Lys Asp Cys
     Val Gly Pro Glu Val Glu Lys Ala Cys Ala Asn Pro Ala Ala Gly Ser
     Val Ile Leu Leu Glu Asn Leu Arg Phe His Val Glu Glu Gly Lys
     Gly Lys Asp Ala Ser Gly Asn Lys Val Lys Ala Glu Pro Ala Lys Ile
     Glu
```

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<211> 148
     <212> PRT
     <213> Homo sapien
Met Ala Thr Leu Lys Glu Lys Leu Ile Ala Pro Val Ala Glu Glu Glu
Ala Thr Val Pro Asn Asn Lys Ile Thr Val Val Gly Val Gly Gln Val
Gly Met Ala Cys Ala Ile Ser Ile Leu Gly Lys Ser Leu Ala Asp Glu
Leu Ala Leu Val Asp Val Leu Glu Asp Lys Leu Lys Gly Glu Met Met
Asp Leu Gln His Gly Ser Leu Phe Leu Gln Thr Pro Lys Ile Val Ala
 Asp Lys Asp Tyr Ser Val Thr Ala Asn Ser Lys Ile Val Val Thr
 Ala Gly Val Arg Gln Gln Glu Gly Glu Ser Arg Leu Asn Leu Val Gln
 Arg Asn Val Asn Val Phe Lys Phe Ile Ile Pro Gln Ile Val Lys Tyr
 Ser Pro Asp Cys Ile Ile Ile Val Val Ser Asn Pro Val Asp Ile Leu
                         135
  Thr Tyr Val Thr
  145
        <210> 216
        <211> 527
        <212> PRT
        <213> Homo sapien
  Gln Arg Ala Pro Gly Ile Glu Glu Lys Ala Ala Glu Asn Gly Ala Leu
  Gly Ser Pro Glu Arg Glu Glu Lys Val Leu Glu Asn Gly Glu Leu Thr
   Pro Pro Arg Arg Glu Glu Lys Ala Leu Glu Asn Gly Glu Leu Arg Ser
   Pro Glu Ala Gly Glu Lys Val Leu Val Asn Gly Gly Leu Thr Pro Pro
   Lys Ser Glu Asp Lys Val Ser Glu Asn Gly Gly Leu Arg Phe Pro Arg
   Asn Thr Glu Arg Pro Pro Glu Thr Gly Pro Trp Arg Ala Pro Gly Pro
    Trp Glu Lys Thr Pro Glu Ser Trp Gly Pro Ala Pro Thr Ile Gly Glu
    Pro Ala Pro Glu Thr Ser Leu Glu Arg Ala Pro Ala Pro Ser Ala Val
    Val Ser Ser Arg Asn Gly Gly Glu Thr Ala Pro Gly Pro Leu Gly Pro
    Ala Pro Lys Asn Gly Thr Leu Glu Pro Gly Thr Glu Arg Arg Ala Pro
    Glu Thr Gly Gly Ala Pro Arg Ala Pro Gly Ala Gly Arg Leu Asp Leu
```

Gly Ser Gly Gly Arg Ala Pro Val Gly Thr Gly Thr Ala Pro Gly Gly  180  180  180  Gly Pro Gly Ser Gly Val Asp Ala Lys Ala Gly Trp Val Asp Asn Thr  200  Arg Pro Gln Pro				
195 Arg Pro Gln Pro Pro Pro Pro Pro Leu Pro Pro Pro Glu Ala Gln 220				
Arg Pro Gln Pro Pro Pro Pro Leu Pro Pro Pro Pro Glu Ala Gln 220 210 217 218 220 219 210 210 210 210 210 210 210 210 210 210				
210 ZIS Pro Clu Val Ala				
210 215 Pro Arg Arg Leu Glu Pro Ala Pro Pro Arg Ala Arg Pro Glu Val Ala Pro Arg Arg Leu Glu Pro Ala Pro Pro Arg Ala Arg Pro Glu Val Ala 235 240				
235 230 235 240 225 230 235 240 Pro Glu Gly Glu Pro Gly Ala Pro Asp Ser Arg Ala Gly Gly Asp Thr 250 255				
Pro Glu Gly Glu Pro Gly Ala Flo Asp Sol Las 255				
Ala Leu Ser Gly Asp Gly Asp Pro Pro Lys Pro Glu Arg Lys Gly Pro 265 270				
260 203  Glu Met Pro Arg Leu Phe Leu Asp Leu Gly Pro Pro Gln Gly Asn Ser 280 285				
275 280 205 Glu Gln Ile Lys Ala Arg Leu Ser Arg Leu Ser Leu Ala Leu Pro Pro 300				
295 300 290 \ 295 Cly Pro Arg Arg Pro Pro Trp				
290 Leu Thr Leu Thr Pro Phe Pro Gly Pro Gly Pro Arg Arg Pro Pro Trp 320 315 316				
305 310 310 Glu Gly Ala Asp Ala Gly Ala Gly Gly Glu Ala Gly Gly Glu Gly Ala Gly Glu Gly Ala Gly 335				
Ala Pro Gly Pro Ala Glu Glu Asp Gly Glu Asp Glu Asp Glu Asp Glu				
345 340 Glu Glu Asp Glu Glu Ala Ala Ala Pro Gly Ala Ala Ala Gly Pro Arg 360 365				
360 365 360 365 360 Yal Pro Val Val Ser Ser				
355 Gly Pro Gly Arg Ala Arg Ala Pro Val Pro Val Val Val Ser Ser 370 375 370 370 370 370				
370 Ala Asp Ala Asp Ala Arg Pro Leu Arg Gly Leu Leu Lys Ser Pro Ala Asp Ala Asp Ala Arg Pro Leu Arg Gly Leu Leu Lys Ser Pro 395 400				
390 395 Arg Lys Arg Lys Arg Lys				
385 Arg Gly Ala Asp Glu Pro Glu Asp Ser Glu Leu Glu Arg Lys Arg Lys 415 405 407				
Met Val Ser Phe His Gly Asp Val Thr Val Tyr Leu Phe Asp GII GIU				
Thr Pro Thr Asn Glu Leu Ser Val Gln Ala Pro Pro Glu Gly ASP 1112				
And Dro Ser Thr Pro Pro Ala Pro Pro Thr Pro Pro His Pro Ala Thr				
450 450 450 450 Asp Gly Phe Pro Ser Asn Asp Ser Gly Phe Gly Gly Ser Phe 475 480				
Pro Gly Asp Gly File F10 Set 1351 1 475 480 475 465 470 Pro Pro Leu				
465 Glu Trp Ala Glu Asp Phe Pro Leu Leu Pro Pro Pro Gly Pro Pro Leu 495 485 486 470 497 497 498				
485  Cys Phe Ser Arg Phe Ser Val Ser Pro Ala Leu Glu Thr Pro Gly Pro 505  505				
500 503 Pro Ala Arg Ala Pro Asp Ala Arg Pro Ala Gly Pro Val Glu Asn				
525 515 520 525				
<210> 217				
<211> 466				
<212> DNA <213> Homo sapien				
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gaatggtgcc tgtcctgctg tctctgctgc tgctcctggg tcotgty accaagatgg tcgttactct ctgacctata tctacactgg gctgtccaag catgttgaag				
accaagatgg tegttactet etgacetata tetacactgg geographic agatacaaca acgteecege gttteaggee ettggeteae teaatgacet ceagttettt agatacaaca				

gtaaagacag gaagteteag eecatgggae tetggagaca ggtggaagga atggaggatt ggaageagga eageeaactt eagaaggeea gggaggacat etttatggag accetgaaag acategtgga gtattacaae gacagtaaeg ggteteaegt attgeaggga aggtttggtt gtgagatega gaataacaga ageageggag eattetggaa atattaetat gatggaaagg actacattga atteaacaaa gaaateeeag eetgggteee ettega	240 300 360 420 466
<210> 218 <211> 381 <212> DNA <213> Homo sapien	
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<210> 219 <211> 1293 <212> DNA <213> Homo sapien	
gaggggaggc gcatggcgg gatggcgtg gatggcgct gagagcagat gagagggaggc gaggggggct tacatacagt acctgctggt cactgaggggct tacatgctgg aggaggaggtg gaggagggggggggg	60 120 180 240 300 360 420 480 540 600 660 720 780 840 900 960 1020 1080 1140 1200 1260 1293
<210> 220 <211> 983 <212> DNA <213> Homo sapien	

caggttatte tgatectgee geetgettee eetgtaagag tggageeteg aggtgaeet tgeagtggeea aatgeeaat tgeagagetg gggeaaggaa gggeteettg aagaegagg taaaaacaga eaagaagtaa ggetggaetg eetteeteag agageeteeteeteeteeteeteeteeteeteeteeteeteet	60 120 180 240 300 360 420 480 540 600 660 720 780 840 900 960 983
<210> 221 <211> 373 <212> DNA <213> Homo sapien	
<400> 221 cattttatgg gttaatttt tattaaatag caataagata cttttataac tcaataaaat tattcaatga tacattcgga aaataaatgt ataaaatatg aaaaagtact aaaaagcatt tttcagtact tttaggtaag attaatccaa ctaaacacta gcatatgtta tacagtaata ataaggggaa aatacaataa tgttgagaaa gcaaactcaa agcatagatc aatgaaaaaa tgtgagaaatg gacataaatg atttagtatt tttaaagaga gtgaaaaatc attatttat gctttgtgt agcgttagat gaattaaata acatatgcac atatagcttt gcgatacaaa tttccagacc ata	60 120 180 240 300 360 373
<210> 222 <211> 544 <212> DNA <213> Homo sapien	
cagagatgct gctgctacaa aggatcggtg taagcagtta acccaggaaa tgatgacaga gaagaaagaa agcaatgtgg ttataacaag gatgaaagat cgaattggaa cattagaaaa ggaacataat gtattcaaa acaaaataca tgtcagttat caagagactc aacagaagga gatgaagttt cagcaagatt ggaggcagaa ggaggcagag atagctcact tgaagcagga gcaagcagtc aactagaaaa gcaagaacta aacaggaaag ctacagcaag aggaagtca aaggaagatca aaggaagatca aaggaagatca aaggaagatca aaggaagaca gctgagaaa gctgagaaa gctgagaaa gaggagtcaa aaggtgggaa gaagttcaag gctacatcaa aaggaagaaca gcggaacatg aggcagcaca gctaagattta cagagtaaat tgttggccaa aaggtgggaa gaagttcaga gctacatcaa aagaagaaca aaggcagcaca gctaagattta cagagtaaat ttgttggccaa aagaagaaaa aaggcagcaca gctaagattta cagagtaaat ttgttggccaa aagaa	60 120 180 240 300 360 420 480 540

<210> 223 <211> 316 <212> DNA <213> Homo sapien

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cagaccacgt ctgecetege cgetetagee ctgegecea geeggeege cetaggatga attacatgga ceggetege ceggetege ceggetegeg cetaggatga aggaagatga aggegaagaa aggegaagatga aggegaagatga aggegaagatga agacgaagatg gatgeteadt tagagaacet cacaagaaga attatgaag cacacagaatgaattggaagaagatggaagaagaagaagaagaag	60 120 180 240 300 360 420 480 540 600 660 720 780 840 900 960 1020 1080 1140 1260 1320 1380 1440 1500 1560 1583
<pre>&lt;210&gt; 225 &lt;211&gt; 491 &lt;212&gt; DNA &lt;213&gt; Homo sapien  &lt;400&gt; 225  gaacaacatc atcttgaatc actagataga ctcttgacgg aaagcaaagg ggaaatgaaa aaggaaaata tgaagaaaga tgaagcttta aaagcattac agaaccaagt atctgaagaa acaatcaagg ttaggcaact agattcagca ttggaaatt gtaaggaaga acttgtcttg catttgaatc aattggaagg aaataaggaa aagtttgaaa aacagttaaa gaagaaatct gaagaggtat attgtttaca gaaagagcta aagataaaaa atcacagtct tcaagagact</pre>	60 120 180 240 300

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tgtgctgagc ctgctggata actacctgat caagaattgc agcgagaccc agtacgagag 540
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<211> 536
<212> DNA
<213> Homo sapiens
<400> 274
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gatggataat gctgactcaa gtcctgtggt agataagaga gaggttattg atttgcttaa 180
acctgaccaa gtagaaggga tccagaaatc tgggactaaa aaactgaaga ccgaaactga 240
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agatgettta aagaagteaa gtaagggaga attgaetaeg ettataeate agetteaaga 420
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 <211> 494
 <212> DNA
 <213> Homo sapiens
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 <222> (379)
 <223> n=A,T,C or G
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caccatcgag aatgtcaagg caaagatcca agataaggaa ggcatccctc ctgaccagca 180
gaggctgatc tttgctggaa aacagctgga agatgggcgc accctgtctg actacaacat 240
ccagaaagag tccaccctgc acctggtgct ccgtctcaga ggtgggatgc aaatcttcgt 300
gaagacactc actggcaaga ccatcaccct tgaggtggag cccagtgaca ccatcgagaa 360
cgtcaaagca aagatccang acaaggaagg cattcctcct gaccagcaga ggttgatctt 420
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<210> 276
<211> 484
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<213> Homo sapiens
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 gctcatcata caccagagga cacacacagg agaaaagcct tatgtctgcg gagagtgtgg 180
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 tgggctggca gctgagttgc agcagcagca ggctgagtac gaggacctta tgggacagaa 300
  agatgacete aacteecage tecaggagte attaegggee aatagtegae tgetggaaca 360
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  gagtgcggag aagcggaagg ccatgcttgg atgagctagc aatggaaacg ctgcaagaga 480
  agtcccacac aaggaagagc ttgggagcag ttc
  <210> 278
  <211> 471
  <212> DNA
  <213> Homo sapiens
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  <400> 278
  caaagaggaa ctcgagcgga ccaacaaaat gctcaaagcc gaaatggaag acctggtcag 120
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  ggatctccaa gcccgggacg agcagaatga ggagaagagg aggcaactgc agagacagct 360
   tcacgagtat gagacggaac tggaagacga gcgaaagcaa cgtgccctgg cagctgcagc 420
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471
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<210> 279
<211> 497
<212> DNA
<213> Homo sapiens
<220>
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<223> n=A,T,C or G
<221> misc_feature
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<223> n=A,T,C or G
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tcagatcact gtcaatggga ccgttctcag ctccagtgga accaggtttg ctgtgaactt 180
teagactgge tteagtggaa atgacattge ettecaette aacceteggt ttgaagatgg 240
agggtacgtg gtgtgcaaca cgaggcagaa cggaagctgg gggcccgagg agaggaagac 300
 acacatgeet ttecagaagg ggatgeeett tgacetetge tteetggtge agageteaga 360
 tttcaaggtg atggtgaacg ggatcctctt cgtgcagtac ttccaccgcg tgcccttcca 420
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 <223> n=A,T,C or G
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 gacagtgggc gcaggccagt gtgtgtgcgc accgtgcgcg agccgaagca gggcgaggca 180
  ttgcctcacc tgggaagcac aaggggtcag ggagttccct ttccgagtca aagaaagggg 240
  tgacggacgc acctggaaaa tcgggtcact cccacccgaa tattgtgctt ttcagaccgg 300
  cttaagaaac ggcgcaccac gagactatat cccacacctg gctcagaggg tcctacgccc 360
  acggaatete getgattget ageacageag tettagatea aactgeaagg ggggeaacga 420
  ggctggggga ggggggcccg ccattgccca ngcttgctta ggtaaacaaa gcagccggga 480
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  cctc
  <210> 281
  <211> 527
  <212> DNA
  <213> Homo sapiens
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<222> (456)
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ataactacac tetetecaag acagagttee taagetteat gaatacagaa etagetgeet 180
tcacaaagaa ccagaaggac cctggtgtcc ttgaccgcat gatgaagaaa ctggacacca 240
acagtgatgg tcagctagat ttctcagaat ttcttaatct gattggtggc ctagctatgg 300
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geeetggeet teaaacceae eccettteet teeageettt etgteateat etceaeagee 420
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<210> 282
<211> 514
 <212> DNA
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 <220>
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 ggtggtcgtg tgtggccagg cgtctgtggg caaaacttca atcctggagc agcttctgta 180
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 agatggggcc gaactgcccc gacactgctt ctcttgcact gatggctacg tcctggtcta 360
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  <210> 283
  <211> 484
  <212> DNA
  <213> Homo sapiens
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  ggaccgcgcc gggaagagca cgcagagccg caagctggtg gaagcgctgt gcgccgcggg 120
  ccaccgcgcc gaactgctcc ggttcccgga aagatcaact gaaatcggca aacttctgag 180
  ttcctacttg caaaagaaaa gtgacgtgga ggatcactcg gtgcacctgc ttttttctgc 240
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  cgtggacaga tacgcatttt ctggtgtggc cttcaccggt gccaaggaga atttttccct 360
  agactggtgt aaacagccag acgtgggcct tcccaaaccc gacctggtcc tgttcctcca 420
   gttacagctg gcggatgctg ccaagcgggg agcgtttggc catgagcgct atgagaacgg 480
   ggct
   <210> 284
   <211> 514
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<212> DNA
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atgagaatta taaagaccca accaacttgc agggaaagct tcagaagcat caagcatttg 360
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gccagtggga attacttttg gagaagatgc gaga
<210> 285
<211> 383
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 cagatgatag gaatcccatc tcatttagaa tggatgacaa aggagaagtg aagtgcatta 300
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 <211> 943
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  cgtggggctg gggcggcgag gctggcggtc cggcctggcc gcgactctgc ccttctttcc 240
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  teeggetgag cacegecaac acetactect accacaaagt ggaettgeee tteeaggagt 360
  atgtggagca gctgctgcac ccccaggacc ccacctccct gggcaatggt gaggcagccc 420
  taggcggcgg tagggggtgg ggacgcttgg agtctccagg tgccaggatc cctgtccccg 480
  ccgtctctgt tggcagacac cctgtacttc ttcggggaca acaacttcac cgagtgggcc 540
  tetetette ggeactaete eccacecca tttggeetge tgggaacege tecagettae 600
  agetttggaa tegeaggage tggetegggg gtgeeettee actggeatgg accegggtae 660
  tcagaagtga tctacggtcg taagcgctgg ttcctttacc cacctgagaa gacgccagag 720
  ttccacccca acaagaccac actggcctgg ctccgggaca catacccagc cctgccaccg 780
  totgcacggc cootggagtg taccatecgg gotggtgagg tgctgtactt coocgaccgc 840
  tggtggcatg ctacgctcaa ccttgacacc agcgtcttca tctccacctt cctcggctag 900
  ccaaaacagc tggcaggact gccggtcaca caccagcacg tcc
   <210> 287
   <211> 1143
   <212> DNA
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<213> Homo sapiens
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gaagatgaac aggagaggcc cttggccctc tgtgaaccag gtgtcaatcc cgaggaacaa 180
ctgattataa tccaaagtcg tctggatcag agtatggagg agaatcagga cttaaagaag 240
gaactgctga aatgtaaaca agaagccaga aacttacagg ggataaagga tgccttgcag 300
cagagattga ctcagcagga cacatctgtt cttcagctca aacaagagct actgagggca 360
aatatggaca aagatgagct gcacaaccag aatgtggatc tgcagaggaa gctagatgag 420
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cagcaccagg ccaagttaga agaagcactc cggaaactct ctgatgtcag ttaccaccag 540
gtggatctag agcgagagct agaacacaaa gatgtcctct tggctcactg tatgaaaaga 600
gaggcagatg aggcgaccaa ctacaacagt cacaactctc aaagcaatgg ttttctcctt 660
ccaacggcag gaaaaggagc tacttcagtc agcaacagag ggaccagcga cctgcagctt 720
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 gtcaatatac caaagaggtt ggaggaggtg acgttaaagg attttaaagc agctattgat 1080
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 <211> 881
 <212> DNA
 <213> Homo sapiens
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  tgagttaatg ggccgggacc gaaacctagc cccggacgag aagcgcagca acgtgcggtg 180
  ggaccacgag agcgtttgta aatattatct ctgtggtttt tgtcctgcgg aattgttcac 240
  aaatacacgt tetgatettg gteegtgtga aaaaatteat gatgaaaate tacgaaaaca 300
  gtatgagaag agetetegtt teatgaaagt tggetatgag agagattttt tgegataett 360
  acagagetta ettgeagaag tagaacgtag gateagaega ggeeatgete gtttggeatt 420
  atctcaaaac cagcagtctt ctggggccgc tggcccaaca ggcaaaaatg aagaaaaat 480
  tcaggttcta acagacaaaa ttgatgtact tctgcaacag attgaagaat tagggtctga 540
  aggaaaagta gaagaagccc aggggatgat gaaattagtt gagcaattaa aagaagagag 600
  agaactgcta aggtccacaa cgtcgacaat tgaaagcttt gctgcacaag aaaaacaaat 660
  ggaagtttgt gaagtatgtg gagccttttt aatagtagga gatgcccagt cccgggtaga 720
  tgaccatttg atgggaaaac aacacatggg ctatgccaaa attaaagcta ctgtagaaga 780
  attaaaagaa aagttaagga aaagaaccga agaacctgat cgtgatgagc gtctaaaaaa 840
  ggagaagcaa gaaagagaaa aaaaaaaaaa aaaaactcga g
   <210> 289
   <211> 987
   <212> DNA
   <213> Homo sapiens
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<210> 292 <211> 511

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tcacagaaca agacaatggt taaaaaccag aacagatgcc cagaaggggg taccatggcc 300
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agtecticag gitgitetig atgatgaeat eggigaegge gicaaacaeg aacigeaegt 600
tettggtgte ggtggegeae gtgaagtgeg tgtagatete ettggtgtet ttgegettat 660
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tatactcagg gaagcagatg gtcaggggac tgtgtgtgat cttctcctca aacaggtcct 780
tettgttgag gaagaggatg atggacgtgt etgtgaacca ettgttgttg cagatgetat 840
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        <210> 290
        <211> 300
        <212> DNA
        <213> Homo sapien
        <220>
        <221> misc feature
        <222> (1)...(300)
        <223> n = A, T, C or G
        <400> 290
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                                                                          60
  tcatcgtggc tgcatccaac ctccgggcag aaaactatga cattccttct gcagaccggc
                                                                         120
  acaagagcaa getgattgca gggaagatca teecageeat tgecaegaee acageageeg
                                                                         180
  tggttggcct tgtgtgtctg gagctgtaca aggttgtgca ggggcaccga cancttgact
                                                                         240
  cctacangaa tgggtgcctc aacttgagcc ctgcctttct ttggtttctc tgaacccett
                                                                          300
        <210> 291
        <211> 352
         <212> DNA
         <213> Homo sapien
         <220>
         <221> misc_feature
         <222> (1)...(352)
         <223> n = A, T, C or G
         <400> 291
   aaccaagctg ccaccggggg tggatcggat gcggcttgag aggcatctgt ctgccgagga
                                                                           60
                                                                          120
   cttctcaagg gtatttgcca tgtcccctga agagtttggc aagctggctc tgtggaagcg
   gaatgagete aagaagaagg eetetetett etgatggeee eeacetgete egggaeggee
                                                                          180
   cccttacccc tgctgcttca gggtttttcc ccggcgggtt gggaggggca ggaggtgggg
                                                                          240
                                                                          300
   tggaaatngg gtgggcncct ttcctcaggt agagnggggg gccaaaacct ctgcngtccc
                                                                          352
   cggagngage tatggaettt etteceete acaaggntgg gggeeteetg et
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<212> DNA
     <213> Homo sapien
     <220>
     <221> misc_feature
     <222> (1)...(511)
     <223> n = A, T, C or G
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ctatgccagg cgcatctcag ctaatccaaa agtaaatgag aaacttagaa aaagattgcc
                                                                        120
aattccaaat caacatattt agagaaaatt ggaaaaggag aagcttacta cagctttatt
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tgaggacttt ttaaagaacg ctgggttcta tctgtgagct gcaaatcttg gagcaaaaac
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cagagacatt gccagagcaa acaagaacag aaatacaaat ggagaactgg tcaaaagaca
                                                                        300
taacccacag ttatcttgaa caagaaacta cggggataaa taaaagtacg canccagatg
                                                                        360
agcaactgac tatgaattct gagaaaagta tgcatcggaa atccactgaa ttagntaatg
                                                                        420
aaataacatg ngagaacaca gaatggccag gggcagagat caacgaattt tcanatcatc
                                                                        480
                                                                        511
agttettate cagatgatga gtetgtttae t
      <210> 293
      <211> 526
       <212> DNA
       <213> Homo sapien
       <220>
       <221> misc_feature
       <222> (1)...(526)
       <223> n = A, T, C or G
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                                                                          60
 acacggtacc accetgetet cetacttete aaacceacat ceaccacea gacaggaggg
                                                                         120
 tgcanacccc acaggaaatt acctcccgga gcactgactg atatttttcc ttaaaacaaa
                                                                         180
 aaaatggctg tctcagacta ataacagaac atcttaagag ctataccagc tattacagcc
                                                                         240
 tggtaatana agcagctttc taanaattcc caagtttata anaggcccaa naaatgcatt
                                                                         300
 tattctgttg tctattaagc ctccatgaca aggagaaagt tatgagtaaa tccttggttc
                                                                          360
                                                                          420
 atcaggagtt aagagctgtg ngcctcatga ggagttaana gctgtgtgca taagcaggtt
 caagaaacaa actcctgttt gtttgcctct ttgatggttc aaaaacattc agctgctttc
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                                                                          526
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   aaggacagtt ctttttgttt tgtttctaat gtcggaagaa aaagaaagag atatattaaa
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actatcattg atgccccagg acacagagac tttatcaaaa acatgattac agggacatct caggctgact gtgctgtcct gattgttgct gctggtgttg gtgaatttga agctggtatc tccaagaatg ggcaggaccc gagagcatgc ccttctggct tacacactgg gtgtgaaaca actaattgtc ggtgttaaca aaatggatt	360 420 480 509
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                                                                        441
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 ctggcgacga aagcagcttc gagctcatct gcggcggtgc caccctcccg ccccggaggc
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 ccggcccttc atatgcggca actgtggccg gagctttgcc cagtgggacc agctagttgc
                                                                         300
                                                                         360
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                                                                         420
 ccggcccagg ggccgccccg cggtgaccgc cccccggccc ggtggagatg ccgtcgaccg
 ccccttccag tgtgcctgtt gtggcaagcg cttccggcac aagcccaact tgatcgctca
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                                                                         491
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  cctgtagaac cagatgttgc ttcaggagat gatacactct gcgttggctt ttcatttctc
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  tggtttggtg tagaaattat aagcctgtct tgccccctga cacttatttc tgttttgtta
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  ccaattccct ttgttgaata aacaaattga tcgataaatt tcccatcccc tgtagcattc
                                                                          360
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ccaggctatt ttaacaggcg gnggctcttc ctctttccgc acttgtgtgc cgcctctggc
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 acataatcta ctatatttga aggactcaaa caaatacatg tttggctgtg aattctgtac
                                                                      180
 tctcaccaaa acagagataa aaatccacct aaaatacact ttccttcatt tagtgcttgt
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 ggganaaggt caagtattgc actttaaaat tactttcatc taacatttgc cccaactttc
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 cccctgaatt cactatatgt tttcagcaaa catgatttta taaattttaa gtataaaagc
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                                                                      420
 aaacggcata tttacttaca aaattganag ataggggcat ccagctgagg tacatttcct
                                                                      480
 cccttggcgt tgagtttctg gacttgggtc gggggcacag gcttgtgtga ctgccccgtg
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                                                                      571
 geocgataca tggcetggae eccaggatge g
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  tcaggtcacc actggatatc agttgcctgt atataattat caggcatttc ctgcttatcc
                                                                       120
  aagttcacca tttcaggtca ccactggata tcagttgcct gtatataatt atcaggcatt
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  tectgettat ecaagtteae cattteaggt caccactgga tateagttge etgtatataa
                                                                       240
  ttatcaggca tttcctgctt atccaagttc accatttcag gtcaccactg gatatcagtt
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  gcctgtatat aattatcagg catttcctgc ttatccaagt tcaccatttc aggtcaccac
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  tggatatcag ttgcctgtat ataattatca ggcatttcct gcttatccaa gttcaccatt
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  teaggteace actggatate agttgeetgt atataattat caggeattte etgettatee
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  aaattcagca gttcaggtca ccactggata tcagttccat gtatacaatt accagatgcc
                                                                        540
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                                                                        420
ggaatcctct agtcaggtgg agcctggaac agacaggaaa tcaactgggt ttgaaactct
ggtggtgacc tccgaagatg gcatcacaaa gatcatgttc aaccggccca aaaagaaaaa
                                                                        480
tgccataaac actgagatgt atcatgaaat tatgcgtgca cttaaagctg ccagcaanga
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                                                                        591
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aagtctcacc tgaaacagtt gacagtgtga ttatgggcaa tgtcctgcag agttcttcag
                                                                        180
atgctatata tttggcaagg catgttggtt tgcgtgtggg aatcccaaag gagaccccag
                                                                        240
ctctcacgat taataggctc tgtggttctg gttttcagtc cattgtgaat ggatgtcagg
                                                                        300
aaatttgtgt taaagaagct gaagttgttt tatgtggagg aaccgaaagc atgagccaag
                                                                        360
ctccctactg tgtcagaaat gtgcgttttg gaaccaagct tggatcagat atcaagctgg
                                                                        420
 aagattettt atgggtatea ttaacagate ageatgteea geteeceatg geaatgaetg
                                                                         480
 cagagaatct tgctgtaaaa cacaaaataa gcagagaaga atgtgacaaa tatgccctgc
                                                                        540
                                                                         591
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       <211> 591
       <212> DNA
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       <220>
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       <223> n = A, T, C \text{ or } G
       <400> 320
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                                                                          60
                                                                         120
 gccggcacca tgtcgaggca ggcgaaccgt ggcaccgaga gcaagaaaat gagctctgag
 ctcttcaccc tgacctatgg tgccctggtc acccagctat gtaaggacta tgaaaatgat
                                                                         180
                                                                         240
 gaagatgtga ataaacagct ggacaaaatg ggctttaaca ttggagtccg gctgattgaa
 gatttcttgg ctcggtcaaa tgttgggagg tgccatgact ttcgggaaac tgcggatgtc
                                                                         300
 attgccaagg tggcgttcaa gatgtacttg ggcatcactc caagcattac taattggagc
                                                                         360
 ccagctggtg atgaattctc cctcattttg gaaaataacc ccttggtgga ctttgtggaa
                                                                         420
```

		•			
actttaa	ata accactcatc ccttatttat t aga tggtccagat ggctngngga g gng tgacagaaat ccggatgaga t	ggcccaagtt	tgtccaggac	accetnaaag	480 540 591
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ctgcttg gtgcago agtttgc tgcggct	(400> 321) ggct ccacacgtgg gccgccgtag ( cagc cacattgaag gatagagtgg ( ctgc tgaaaatgaa gggaagtctg ( caag tccagagcct ggcagggagg ( gngg anaggagggg	cagcagaggc qqqqaggtct	caaggategt	gagilgalgg gctgaggggg	60 120 180 240 260
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	<210> 323 <211> 492 <212> DNA <213> Homo sapien				
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	<400> 323				

cetqtetece agecqtacca gegagggete ggecggeage geegggetgg ggageggegg egageggggg ggageggggg egagegggggggg	60 120 180 240 300 360 420 480 492
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<210> 325 <211> 532 <212> DNA <213> Homo sapien	
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qaggagacag gacagaggt ctggagagge aggaggacac cgagttccce gtgttggct cagggagacag gacagaggt ctggagagge aggaggacac cgagttccce gtgttgcaac caggtcctg tgcttgcga gccgtccgge ggctgggate gagccccgac aatgggcaac cggcaggage ggccgtcaga gactatcgac cgcgagcgga aacgcctggt caccgggcac caggagcgac cattggatge cattggatge cattggatge caggagcgca caggagcgca cggagcgcac cggaccccgc ttgggactge cagcacgtgg gtccggagcgcaccggagcaccacagcaccacagcaccacagcaccac	60 120 180 240 300 360 420 480 532
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<sup>&</sup>lt;210> 326 <211> 322 <212> DNA

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<210> 327 <211> 387 <212> DNA <213> Homo sapien	
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<210> 328 <211> 502 <212> DNA <213> Homo sapien	
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<210> 329 <211> 463 <212> DNA <213> Homo sapien	·

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     <222> (1) ... (463)
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aagtoottta tacaaaataa ggacaatttg taaaganaat ccactgtoat gttttgoott
                                                                        120
gtcaagtcaa aactcaaata gcttgttttg gtaaaattat tccagaaaca taatccagac
                                                                        180
aaaatcaata acgtcatcag cttcctaacc atgtttaana ggaataactt catgaacatt
                                                                        240
ttgccctgaa ctgaanagtt ctaaatactt gtaaaccttt aggaaaaaat gactgctcgc
                                                                        300
aggcagcttg actggtaaga gggtacacca nagactccgg gtcactcact gtcagaatat
                                                                        360
tettatacat acaatgagte tecaegeetg tacaatgagt gtegtgeaac ataattggag
                                                                        420
                                                                        463
taatggcctc taaaatttta caagtaaact ttattgnggc ccc
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      <211> 500
       <212> DNA
       <213> Homo sapien
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       <221> misc_feature
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                                                                          60
                                                                         120
 caaaaggact ataaaacaaa aacagagaag aaaattcatg gctaaaccag ctgaagaaca
 gcttgatgtg ggacagtcta aagatgaaaa catacataca tcacatatta cccaagacga
                                                                         180
                                                                         240
 atttcaaaga aattcagaca gaaatatgga agagcatgaa gagatgggaa atgattgtgt
                                                                         300
  ttccaaaaaa acagatgcca cctgtgggaa gcaagaaaag tagcactaga aaagataagg
  aagaatctaa aaagaagcgc ttttccagtg agtccaagaa caaacttgtn cctgaagaag
                                                                          360
  tgacttcaac tgtcacgaaa agtcgaanaa tttccangcg tccatctgat tggtgggtgg
                                                                          420
  taaaancaga ggagagtoot gtttatagca attottoagt aagaaatgaa ttaccaantg
                                                                          480
                                                                          500
  catcacaatn ntgcccggaa
        <210> 331
        <211> 494
        <212> DNA
        <213> Homo sapien
        <220>
        <221> misc_feature
         <222> (1)...(494)
         <223> n = A,T,C or G
                                                                            60
   tetetetete teteaaaatt acagtgttea ttgteattga eeteageage aaatttgaet
   tgaattcact taggatcgca ggaatcaggg gaaagtgatt ttaaaggtgg tttctccagc
                                                                           120
   acattttaag aaaagggacc aaaagttatt ttagcttcct caatagattg catgttgctt
                                                                           180
   attaggataa taaattaata ttaaatgcaa tatatgtott gnotttatta tggcatotat
                                                                           240
   ttaggagttg ttcaaatcac tgcagtaggg ctctgcaaat aaaataatgn aacctattat
                                                                           300
   catggatcta atgnactgna actttatcag tgaaaggnaa aatctcaaat aacaagtaca
                                                                           360
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threatat agagattigt agaggaaaa titticcata gatticates	20 180 194
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<210> 333 <211> 499 <212> DNA <213> Homo sapien	
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tota atgaaattat gcgtgcactt aaagctgcca gcaaggatgac	480 540 561
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aagetggtea tggetggga gaccaccaac teeegeggee ageggetgee ceagaaggga gacgetggee tgeeaggget teageggeat teeetgeggee atteeteetg gtteaaaaac teeetggtgg tteetteeteetg gagaatgtea gaacetteteetgeeteeteetgeeteeteetgeeteeteetgeeteet	60 120 180 240 300 360 420 480 540 551
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<210> 337 <211> 422 <212> DNA <213> Homo sapien	
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<400> 337 gcagcaggaa cagttacagc agcagcagca acagcagctg ttgcaacagc agcaggaaca	a 60

attgcagcag caacaactgc agcctcctcc cctggagccc gaggaggagg aagaggtgga attgcagcag caacaactgc agcctggggtc agagcaggag ctggagcagc agctcaaact	120 180 240 300 360 420 422
<210> 338 <211> 601 <212> DNA <213> Homo sapien	
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catettacga acgetetatg atgetetatg ageggtetat gatgteecet atggetgaac etactatgat gteageetae gagegeteta tgatgteage etacagagge tectatgatg eccetatgge tgagegetet atgatgteage etacatgatg etecatgatg etacatgatg etatgatget atggetgate atggetgate gteategate getgetgate etatgatgte getgetatet etatactgetg atggetetatgat etatgatget etatacetet gatgteatet etatacetget gatgteatat etatactgetg eccetagaggetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgetetatgatgatgatgetetatgatgetetatgatgetetatgatgetetatgatgatgatgetetatgatgatgatgetetatgatgatgatgetetatgatgatgatgetetatgatgatgatgetetatgatgatgatgatgetetatgatgatgatgatgatgatgatgatgatgatgatgatg	60 120 180 240 300 360 420 480 540 600
<210> 339 <211> 440 <212> DNA <213> Homo sapien	
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agagggagga ggcccaactg gtgatgctgc tgctgctgct gctgccgccg ccgccgcctc agagggagga ggcccaactg ggctggaagg gtggttccta ttcgcaccat cgccaaccag gactaggagga aaaaaaaaaa	60 120 180 240 300 360 420 440
<210> 340 <211> 450 <212> DNA	

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<213> Homo sapien
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     <222> (1)...(450)
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                                                                        60
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aggatteete aggeegacea gtggaagtet teaaacaaga geetggtgga ggetetgggg
ctggaagccg agggtgcagt tectgagaca cagactttga ccggatggag taaggggtte
                                                                       180
attggcatgc acagggaaat gcaagtcaac cccatttcaa agcggatggg gcccatgact
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gtggtcagga tggacgcttc agtccagcca ggcccttttc ggaccctgct ccagtttctt
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tatacgggac aactggatga aaaggaaaag gatttggtgg gcctggctca gatcgcagag
                                                                        360
gtcctcgaga tgttcgattt gaggatgatg gtggaaaaca tcatgaacaa ggaagccttc
                                                                        420
                                                                        450
atgaaccagg agattacgaa nncctttcac
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       <211> 451
       <212> DNA
       <213> Homo sapien
                                                                          60
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 cateccatga ggatetgeec getteccagg aaaggteega ggttaateea geaegtatgg
                                                                         120
 ggccaagtgt aggctcccag caggaactga gagcgccatg tcttccagta acctatcagc
                                                                         180
                                                                         240
 agacaccagt gaacatggaa aagaacccaa gagaggcacc teetgttgtt eeteetttgg
 caaatgctat ttctgcagct ttggtgtccc cagccaccag ccagagcatt gctcctcctg
                                                                         300
                                                                         360
  ttcctttgaa agcccagaca gtaacagact ccatgtttgc agtggccagc aaagatgctg
                                                                         420
  gatgtgtgaa taagagtact catgaattca agccacagag tggagcagag atcaaagaag
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  ggtgtgaaac acataaggtt gccaacacaa g
        <210> 342
        <211> 498
         <212> DNA
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         <221> misc_feature
         <222> (1)...(498)
         <223> n = A, T, C or G
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                                                                            60
                                                                           120
   actgatactc caaacaagaa accaactaaa ggcaaaggta aaaaacatga agcagatgag
   ttgagtggag atgettetgt gggaagatga tgettttate aaggaetgtg aattggagaa
                                                                           180
   tcaagaggca catgagcaag atggaaatga tgaactaaag gactctgaag aatttggtga
                                                                           240
   aaatgaagaa gaaaatgtgc attccaagga gttactctct gcagaagaaa acaagagagc
                                                                           300
   tcatgaatta atagaggcag aaggaataga agatatagaa aaagaggaca tcgaaagtca
                                                                           360
    ggaaattgaa gctcaagaag gtgaagatga tacctttcta acagcccaag atggtgagga
                                                                           420
    agaagaaaat gagaaagata tagcagggtt ctggtgatgg cncacaagaa gtatntaaac
                                                                           480
                                                                           498
    ctcttccttc aaaaaggg
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<211> 491 <212> DNA <213> Homo sapien	
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<210> 345 <211> 498 <212> DNA <213> Homo sapien	
actagtctc gggccatcct ttctgcgcac ccggtgtcgc tgggctgcac cccgggcggg actagtccgcc gggcacggga gggggccaag atgccgatca ataaatcaga gaagccagaa agctgcgata atgtgaaggt tgttgttagg tgccggccc tcaatgagag agagaaatca agtgcgatca aacaggctgt cagtgtggat gagatgaggg gaactatcac tgtacataag actgattctt ccaatgaacc tccaaagaca tttacttttg atactgatt tggaccagag ggctacaatg ggactatttt tgcatatga caaaccggaa caggcaaaac tttaccatg gaaaggtgtc gagctattcc tgaacttaga ggaataattc cccaatttct ttgctcacaa tatttgggcc atatttgc	60 120 180 240 300 360 420 480 498
<210> 346 <211> 427 <212> DNA <213> Homo sapien	

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Tyr Leu Gln Leu Trp Arg Leu Leu Leu Tyr Arg Glu Arg Arg Leu 75 70	
Tyr Gln Ser Leu Cys Leu Phe Leu Cys Leu Leu Trp Ala Ala Leu 95	
Thr Thr Leu Phe Ser Ala Ala Phe Ser Leu Ser Gly Ser Leu Pro 100 105	
Leu Arg Pro Pro Ala His Leu His Phe Phe Pro His Trp Leu Leu 125	
Cys Phe Pro Ser Cys Leu Gln Phe Ser Thr Leu Cys Leu Leu Asn 130 135	
Tyr Leu Ala Glu Val Ile Cys Lys Val Arg Cys Ala Thr Glu Leu 155 145	
Arg His Lys Ile Leu Leu His Leu Gly Phe Ile Met Ala Ser Leu 175	
Phe Leu Val Val Asn Leu Thr Cys Ala Met Leu Val His Gly As 190	
Pro Glu Asn Gln Leu Lys Trp Thr Val Phe Val Arg Ala Leu Il 200 205	
Asp Ser Leu Phe Ile Leu Cys Ala Ile Ser Leu Val Cys Tyr Il 210 215	
Lys Ile Thr Lys Met Ser Ser Ala Asn Val Tyr Leu Glu Ser Ly 235 230	
Met Ser Leu Cys Gln Thr Val Ile Val Gly Ser Val Val Ile L $^{\prime}$	∍u Leu

255 250 245 Tyr Ser Ser Arg Ala Cys Tyr Asn Leu Val Val Val Thr Ile Ser Gln 260 Asp Thr Leu Glu Ser Pro Phe Asn Tyr Gly Trp Asp Asn Leu Ser Asp Lys Ala His Val Glu Asp Ile Ser Gly Glu Glu Tyr Ile Val Phe Gly 275 Met Val Leu Phe Leu Trp Glu His Val Pro Ala Trp Ser Val Val Leu 290 310 Phe Phe Arg Ala Gln Arg Leu Asn Gln Asn Leu Ala Pro Ala Gly Met 325 Ile Asn Ser His Ser Tyr Ser Ser Arg Ala Tyr Phe Phe Asp Asn Pro 340 Arg Arg Tyr Asp Ser Asp Asp Asp Leu Pro Arg Leu Gly Ser Ser Arg Glu Gly Ser Leu Pro Asn Ser Gln Ser Leu Gly Trp Tyr Gly Thr Met 355 Thr Gly Cys Gly Ser Ser Ser Tyr Thr Val Thr Pro His Leu Asn Gly 370 390 Pro Met Thr Asp Thr Ala Pro Leu Leu Phe Thr Cys Ser Asn Leu Asp 405 Leu Asn Asn His His Ser Leu Tyr Val Thr Pro Gln Asn 420 <210> 392 <211> 1584 <212> DNA <213> Homo sapiens ggaagactgg agcetttgcg geggegetge eceteceetg gteecegega geteggaggg 60 cccggctggt gctgcggggg ccccggggagg ttgaaaacta agcatgggga agagctgcaa 120 ggtggtcgtg tgtggccagg cgtctgtggg caaaacttca atcctggagc agcttctgta 180 tgggaaccat gtagtgggtt cggagatgat cgagacgcag gaggacatet acgtgggete 240 cattgagaca gaccgggggg tgcgagagca ggtgcgtttc tatgacaccc ggggggctccg 300 agatggggcc gaactgccc gacactgctt ctettgcact gatggctacg tcctggtcta 360 tagcacagat agcagagagt cttttcagcg tgtggagctg ctcaagaagg agattgacaa 420 atccaaggac aagaaggagg tcaccatcgt ggtccttggc aacaagtgtg acttacagga 480 gcagcggcgt gtagacccag atgtggctca gcactgggcc aagtcagaga aggtgaagct 540 gtgggaggtg tcagtggcgg accggcgctc cctcctggag ccctttgtct acttggccag 600 caagatgacg caaccccaga gcaagtctgc cttccccctc agccggaaga acaagggcag 660 cggctccttg gatggctgaa gagctgccgt tcctctttca cgatcccagc cccatttcag 720

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Thr Asp Arg Gly Val Arg Glu Gln Val Arg Phe Tyr Asp Thr Arg Gly 50 50
Leu Arg Asp Gly Ala Glu Leu Pro Arg His Cys Phe Ser Cys Thr Asp 70 80
65  Gly Tyr Val Leu Val Tyr Ser Thr Asp Ser Arg Glu Ser Phe Gln Arg 95 85
Val Glu Leu Leu Lys Lys Glu Ile Asp Lys Ser Lys Asp Lys Lys Glu 100 105
Val Thr Ile Val Val Leu Gly Asn Lys Cys Asp Leu Gln Glu Gln Arg 120
Arg Val Asp Pro Asp Val Ala Gln His Trp Ala Lys Ser Glu Lys Val
Lys Leu Trp Glu Val Ser Val Ala Asp Arg Arg Ser Leu Leu Glu Pro 150 155 160
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115

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Gln Gln Met His Pro Thr Val Val Ile Ser Ala Tyr Arg Lys Ala Leu Asp Asp Met Ile Ser Thr Leu Lys Lys Ile Ser Ile Pro Val Asp Ile 130 150 Ser Asp Ser Asp Met Met Leu Asn Ile Ile Asn Ser Ser Ile Thr Thr Lys Ala Ile Ser Arg Trp Ser Ser Leu Ala Cys Asn Ile Ala Leu Asp Ala Val Lys Met Val Gln Phe Glu Glu Asn Gly Arg Lys Glu Ile Asp Ile Lys Lys Tyr Ala Arg Val Glu Lys Ile Pro Gly Gly Ile Ile Glu Asp Ser Cys Val Leu Arg Gly Val Met Ile Asn Lys Asp Val Thr His 210 Pro Arg Met Arg Arg Tyr Ile Lys Asn Pro Arg Ile Val Leu Leu Asp Ser Ser Leu Glu Tyr Lys Lys Gly Glu Ser Gln Thr Asp Ile Glu Ile 260 Thr Arg Glu Glu Asp Phe Thr Arg Ile Leu Gln Met Glu Glu Tyr Ile Gln Gln Leu Cys Glu Asp Ile Ile Gln Leu Lys Pro Asp Val Val Ile Thr Glu Lys Gly Ile Ser Asp Leu Ala Gln His Tyr Leu Met Arg 290 310 Ala Asn Ile Thr Ala Ile Arg Arg Val Arg Lys Thr Asp Asn Asn Arg Ile Ala Arg Ala Cys Gly Ala Arg Ile Val Ser Arg Pro Glu Glu Leu Arg Glu Asp Asp Val Gly Thr Gly Ala Gly Leu Leu Glu Ile Lys Lys Ile Gly Asp Glu Tyr Phe Thr Phe Ile Thr Asp Cys Lys Asp Pro Lys Ala Cys Thr Ile Leu Leu Arg Gly Ala Ser Lys Glu Ile Leu Ser Glu Val Glu Arg Asn Leu Gln Asp Ala Met Gln Val Cys Arg Asn Val Leu

405

Leu Asp Pro Gln Leu Val Pro Gly Gly Gly Ala Ser Glu Met Ala Val

Ala His Ala Leu Thr Glu Lys Ser Lys Ala Met Thr Gly Val Glu Gln

Trp Pro Tyr Arg Ala Val Ala Gln Ala Leu Glu Val Ile Pro Arg Thr

Leu Ile Gln Asn Cys Gly Ala Ser Thr Ile Arg Leu Leu Thr Ser Leu 465

Arg Ala Lys His Thr Gln Glu Asn Cys Glu Thr Trp Gly Val Asn Gly 485

Glu Thr Gly Thr Leu Val Asp Met Lys Glu Leu Gly Ile Trp Glu Pro

Leu Ala Val Lys Leu Gln Thr Tyr Lys Thr Ala Val Glu Thr Ala Val

Leu Leu Leu Arg Ile Asp Asp Ile Val Ser Gly His Lys Lys Gly

Asp Asp Gln Ser Arg Gln Gly Gly Ala Pro Asp Ala Gly Gln Glu 550 545

<210> 397

<211> 307

<212> PRT

<213> Homo sapiens

Arg Glu Ser Arg Ser Arg Ala Met Glu Glu Glu Ala Ser Ser Pro Gly

Leu Gly Cys Ser Lys Pro His Leu Glu Lys Leu Thr Leu Gly Ile Thr

Arg Ile Leu Glu Ser Ser Pro Gly Val Thr Glu Val Thr Ile Ile Glu

Lys Pro Pro Ala Glu Arg His Met Ile Ser Ser Trp Glu Gln Lys Asn

Asn Cys Val Met Pro Glu Asp Val Lys Asn Phe Tyr Leu Met Thr Asn

Gly Phe His Met Thr Trp Ser Val Lys Leu Asp Glu His Ile Ile Pro

Leu Gly Ser Met Ala Ile Asn Ser Ile Ser Lys Leu Thr Gln Leu Thr

110 105 100 Gln Ser Ser Met Tyr Ser Leu Pro Asn Ala Pro Thr Leu Ala Asp Leu 115 Glu Asp Asp Thr His Glu Ala Ser Asp Asp Gln Pro Glu Lys Pro His Phe Asp Ser Arg Ser Val Ile Phe Glu Leu Asp Ser Cys Asn Gly Ser 150 Gly Lys Val Cys Leu Val Tyr Lys Ser Gly Lys Pro Ala Leu Ala Glu Asp Thr Glu Ile Trp Phe Leu Asp Arg Ala Leu Tyr Trp His Phe Leu Thr Asp Thr Phe Thr Ala Tyr Tyr Arg Leu Leu Ile Thr His Leu Gly Leu Pro Gln Trp Gln Tyr Ala Phe Thr Ser Tyr Gly Ile Ser Pro Gln Ala Lys Gln Trp Phe Ser Met Tyr Lys Pro Ile Thr Tyr Asn Thr Asn 210 Leu Leu Thr Glu Glu Thr Asp Ser Phe Val Asn Lys Leu Asp Pro Ser 245 Lys Val Phe Lys Ser Lys Asn Lys Ile Val Ile Pro Lys Lys Gly Pro Val Gln Pro Ala Gly Gly Gln Lys Gly Pro Ser Gly Pro Ser Gly Pro Ser Thr Ser Ser Thr Ser Lys Ser Ser Ser Gly Ser Gly Asn Pro 290 Thr Arg Lys 305 <210> 398 <211> 416 <212> DNA <213> Homo sapiens agaattcggc acgaggattg cctatctcca gtgcaacaac catcaagtgt gctgaaagtc 60 ttcagccggt tgctgcagca gtggaagaaa gggctacagg tccagtcttg ataagcaccg 120 ccgactttga ggggcctatg cccagtgcgc ccccagaagc tgaaagtcct cttgcctcaa 180  aggettetgt tteeggtgta gttgttgaaa gtgaaaatga gegagetgge acagteatgg 300

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aagaaaaaga cgggagtggc atcatctcta cgagctcggt ggaagactgt gagggcccag 360
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      aaaagttcag ccccgcgggg cctgtgctgt ncatccgggt ctgccgngat atgatcaccc 120
       geogetecet gggetatgee taegneaact tecancaace ggeogacget gategggett 180
       tggacaccat gaactttgat gtgattnagg gaaanccaat ccttatcntg tnnnaatcat 240
aggnatcctt ctttgacaa
       <210> 400
       <211> 410
       <212> DNA
<213> Homo sapiens
Į.J
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Ħ
        agttaccate acacceggg aggageegea getgeegeag eeggeeeeag teaccateae 120
1:40
        cgcaaccatg agcagcgagg ccgagaccca gcagccgccc gccgccccc cccgccgccc 180
 M
        cogcoctcag cgccgccgac accaagcccg gcactacggg cagcggcgca gggagcggtg 240
 ļ
        gcccgggcgg cctcacatcg gcggcgcctg ccggcgggga caagaaggtc atcgcaacga 300
        aggttttggg aacagtaaaa tggttcaatg taaggaacgg atatggtttc atcaacagga 360
 1.4
        atgacaccaa ggaagatgta tttgtacacc agactgccat aaagaagaat
 1.2
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         <211> 433
         <212> DNA
         <213> Homo sapiens
         <220>
         <221> misc_feature
         <222> (1)...(433)
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          ccacggagec teeggggetg eeggcacagt etteactace gtagaagace tiggetecaa 120
          gatactecte acctgetect tgaatgacag egecacagag gteacaggge accgetgget 180
          gaaggggggc gtggtgctga aggaggacgc gctgcccggc cagaaaacgg agttcaaggt 240
          ggactccgac gaccagtggg gagagtactc ctgcgtcttc ctccccgagc ccatgggcac 300
          ggccaacate cagetecacg ggccteccag agtgaaggee gtgaagtegt cagaacacat 360
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ctgggcctgg tac

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<211> 434
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<213> Homo sapiens
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gaagegtgee teaggeeagg ettttgaget gatteteage eeteggteaa aaggatetgt 180
tccagaattc cccctttccc ctccaaagaa gaaggatctt tccctggagg aaattcagaa 240
gaaattagaa getgeagaag aaagaegeaa gteecatgaa getgaggtet tgaageaget 300
ggctgagaaa cgagagcacg agaaagaagt gcttcagaag gcaatagaag agaacaacaa 360
 cttcagtaaa atggcagaag agaaactgac ccacaaaatg gaagctaata aagagaaccg 420
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  <212> DNA
  <213> Homo sapiens
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  agtgaagcag aagaaaaata ctgacaaaga ccatccgaat actggaaaca aaaaaggatc 180
  ccattcaaat tcaagaaaaa atattgataa gactgctgtg actagtggaa atcatgtatg 240
  tccttgtaaa gaaagcgaaa cgtttgtaca gtttgccaat ccatcacagc ttcagtgcag 300
  tgataatgta aaaattgttt tagacaagaa tcttaaagat tgcactgagc ttgtcttaaa 360
   gcaacttcag gaaatgaaac ctaccgtcag tctgaaaaaa cttgaagtac attcaaatga 420
   tccagatatg tctgt
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   <212> DNA
   <213> Homo sapiens
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    caatgagtaa acatatteet caattetgtg gtgttettgg teacacattt atggagttte 180
    tgaagggcag tggagattac tgccaggcac agcacgacct ctatgcagac aagtgaactg 240
    tagaaactga ttactgctcc accaagaagc ccccataaga gtggttatcc tggacacaga 300
    agtgttgaat tgaaatccac agagcatttt acaagagttc tgacctggat ggggtaaacc 360
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     <211> 435
     <212> DNA
     <213> Homo sapiens
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     gccatgctgc tatccgtgcc gctgctgctc ggcctcctcg gcctggccgt cgccgagcct 120
      gccgtctact tcaaggagca gtttctggac ggagacgggt ggacttcccg ctggatcgaa 180
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tccaaacaca agtcagattt tggcaaattc gttctcagtt ccggcaagtt ctacggtgac 240
     gaggagaaag ataaaggttt gcagacaagc caggatgcac gcttttatgc tctgtcggcc 300
     agtttcgagc ctttcagcaa caaaggccag acgctggtgg tgcagttcac ggtgaaacat 360
     gagcagaaca tcgactgtgg gggcggctat gtgaagctgt ttcctaatag tttggaccag 420
     acagacatgc acgga
      <210> 406
      <211> 424
      <212> DNA
      <213> Homo sapiens
      <220>
      <221> misc_feature
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       gtataggcga tagaaattga aacctggcgc aatagatata gtaccgcaag ggaaagatga 120
       aaaattataa ccaagcataa tatagcaagg actaacccct ataccttctg cataatgaat 180
taactagaaa taactttgca aggagagcca aagctaagac ccccgaaacc agacgagcta 240
       cctaagaaca gctaaaagag cacacccgtc tatgtagcaa aatagtggga agatttatag 300
       gtagaggcga caaacctacc gagcctggtg atagctggtt gtccaagata gaatcttagt 360
       tcaactttaa atttgcccac agaaccctct aaatcccctt gnaaatttaa ctgntagtcc 420
IJ
        aaag
1
        <210> 407
Ei
        <211> 423
        <212> DNA
 M
        <213> Homo sapiens
 į.£
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 ctgcagccat ggctcccggc cagctcgcct tatttagtgt ctctgacaaa accggccttg 120
         tggaatttgc aagaaacctg accgctcttg gtttgaatct ggtcgcttcc ggagggactg 180
         caaaagctct cagggatgct ggtctggcag tcagagatgt ctctgagttg acgggatttc 240
         ctgaaatgtt ggggggacgt gtgaaaactt tgcatcctgc agtccatgct ggaatcctag 300
         ctcgtaatat tccagaagat aatgctgaca tggccagact tgatttcaat cttataagag 360
         ttgttgcctg caatctctat ccctttgtaa agacagtggc ttctccaggt gtaagtgttg 420
         agg
          <210> 408
          <211> 424
          <212> DNA
          <213> Homo sapiens
          gaaaaaaaat agcttactga attctataag atgtgtggga atctcaccta tcaaaaatag 60
          gtaaaaagag cetecaaace tgetttgatt ttatteacet attetttag geeaggaact 120
          aatttacctc tcactatcct gttccctctt gctatcttgt ggagtctcta aagacaaagg 180
          tataaagagc ttttggtagg tgaattaata atcaactaga tggcatttcc aaatgggatt 240
          gcacatactg tggggcaagt cccaagtgaa cttcaaagtg agacgtttat ttgagtaatc 300
           cttccagatt aacaataatc ataatagcag ttaccacttc ctgagtactt tctatatgcc 360
           atgtattgag cttgctcact tctttatgtg gattcttatt taatcttaat accaagatga 420
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424

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<210> 409
<211> 398
<212> DNA
<213> Homo sapiens
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 cgcctcctaa tccctagcca ctatgcgtga gtgcatctcc atccacgttg gccaggctgg 120
 tgtccagatt ggcaatgcct gctgggagct ctactgcctg gaacacggca tccagcccga 180
 tggccagatg ccaagtgaca agaccattgg gggaggagat gactccttca acaccttctt 240
 cagtgagacg ggcgctggca agcacgtgcc ccgggctgng tttgtagact tggaacccac 300
 agtnattgat gaagntcgna ctggcaccta cccgcaggtc ttncaccctg ancanntcat 360
 nacaggcaag gaagatgctg ncaaataact atgcccga
  <210> 410
  <211> 423
  <212> DNA
  <213> Homo sapiens
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  getegetege teteacgetg eceggecage ecgettetet geceggagee atgaatetea 120
  gtagcgccag tagcacggag gaaaaggcag tgacgaccgt gctctggggc tgcgagctca 180
  gtcaggagag gcggacttgg accttcagac cccagctgga ggggaagcag agctgcaggc 240
  tgttgcttca tacgatttgc ttgggggaga aagccaaaga ggagatgcat cgcgtggaga 300
   tectgecece ageaaaceag gaggacaaga agatgeagee ggteaceatt geeteactee 360
   aggecteagt cetececatg gtetecatgg taggagtgea getttetece ceagttactt 420
   tcc
   <210> 411
   <211> 424
   <212> DNA
   <213> Homo sapiens
    <220>
    <221> misc_feature
    <222> (1)...(424)
    <223> n = A, T, C or G
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    ggaccggccg aacgcagagg ttgattcttc accacactga aaccattagg aaaaatcctt 120
    gtggttaaca gcagaggctt cagagtgtaa cctgtactcg ggcctagaaa ttatttaaaa 180
    tggcgactga tacgtctcaa ggtgaactcg tccatcctaa ggcactccca cttatagtag 240
    gagetcaget gatecaegeg gacaagttag gtgagaaggt agaagatage accatgeega 300
    ttcgtcgaac tgtgaattct acccgggaaa ctcctcccaa aagcaagctt gctgaagggg 360
     aggaagaaan gccagaacca gacataagtt cagaggaatc tgtctccact gtagaagaac 420
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424
     aaga
     <210> 412
     <211> 430
     <212> DNA
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     ctacggccgc cececteceg atgtggaggg tatgacetec etcaaggtgg acaacetgae 120
     ctaccgcacc tegecegaca egetgaggeg egtettegag aagtacggge gegteggega 180
      cgtgtacatc ccgcgggatc gctacaccaa ggagtcccgc ggcttcgcct tcgttcgctt 240
      tcacgacaag cgcgacgctg aggacgctat ggatgccatg gacggggccg tgctggacgg 300
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      taggcggcgt
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       <211> 429
       <212> DNA
       <213> Homo sapiens
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1.4
       acaggaaggc ttcggctgcg tggtcaccaa ccgattcgac cagttatttg acgacgaatc 180
W
       ggaccccttc gaggtgctga aggcagcaga gaacaagaaa aaagaagccg gcgggggcgg 240
Ų
       cgttgggggc cctggggcca agagcgcagc tcaggccgcg gcccagacca actccaacgc 300
şi
       ggcaggcaaa cagctgcgca aggagtccca gaaagaccgc aagaacccgc tgcccccag 360
cgttggcgtg gttgacaaga aagaggagac gcagccgccc gtggcgctta agaaagaagg 420
M
[.±
        aataagacg
 ]:.±
        <210> 414
 14
        <211> 429
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        <213> Homo sapiens
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        caggtgaatg gtaatgcggt acagttaaaa gaagaggaag aaccaatgga tacttccagt 120
        gtaactcaca cagaacacta caagacactg atagaggcag gcctcccaca gaaggtggca 180
         gaaagacttg atgaaatatt tcagacagga ttggtagctt atgtcgatct tgatgaaaga 240
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         aaggaaagtg acttatcaca tgttcagaac aaaagtgcat ttttatgtgg agttatgaag 360
         acctacaggc agagagaga acaggggagc aaggtgcaag agtccacaaa gggacctgat 420
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     cgctctcgcc gaggaacaag tcggtcagga agcccgcgcg caacagccat ggcttttaag 120
     gataccggaa aaacacccgt ggagccggag gtggcaattc accgaattcg aatcacccta 180
     acaagccgca acgtaaaatc cttggaaaag gtgtgtgctg acttgataag aggcgcaaaa 240
     gaaaagaatc tcaaagtact ttgagaatca ctacaagaaa aactccttgt ggtgaaggtt 300
      ctaagacgtg ggatcgtttc cagatgagaa ttcacaagcg actcattgac ttgcacagtc 360
      cttctgagat tgttaagcan attacttcca tcantatt
      <210> 416
      <211> 269
      <212> DNA
      <213> Homo sapiens
<220>
       <221> misc_feature
       <222> (1)...(269)
       <223> n = A, T, C or G
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       tecceggegg geagececag getggteece geeteegete tecceaeegg eggggaaage 120
       agctggtgtg ggaggaaagg ctccatecec egececetet etecegetgt tggetggean 180
1
       gatcttttgg cagtcctgtg gnctcnctcc ccgnccggat cctnctgacc ctganattcn 240
Ų
#
       nggtntnacn nnccgtncac gccttgntt
I
        <210> 417
 1,3
        <211> 408
 1=
        <212> DNA
 2
        <213> Homo sapiens
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         ccctggctga gggagtattg ctgaggacct cacaggagcc cacttectcg gaggtggtga 180
         gctatgcccc attcacgctc ttcccctcac tggtccccag tgccctgctg gagcaagcct 240
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         tggagcaaac tctttccagc accatcaaac aggatgactt taccgctcgt ctctttgaca 360
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          <211> 402
          <212> DNA
          <213> Homo sapiens
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          gtegggeege gageggagee ggetgagegg gegeegaget eeegeeatgg eeeggaacae 120
          gctgtcctcg cgcttccgcc gggtggacat cgacgaattt gacgagaaca aatttgtgga 180
          cgagcaggag gaggcggcgg cggcggcggc ggagccaggc ccggacccga gcgaggtgga 240
          cgggctcctg cggcaagggg acatgcttcg ggcattccat gcagccttgc ggaactctcc 300
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```
cgtcaacacc aagaatcaag ctgtgaagga gcgagcccag ggcgtggtgc tgaaagtgct 360
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     <210> 419
     <211> 406
      <212> DNA
     <213> Homo sapiens
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Asp Asp Ser Arg Glu Gly Glu Leu Met Asp Cys Asp Gly Lys Ser Glu

Ser Ser Pro Glu Arg Glu Ala Val Asp Asp Glu Thr Lys Gly Val Glu 310 315
Gly Thr Asp Gly Val Lys Lys Arg Lys Arg Lys Pro Tyr Arg Pro Gly 335
Ile Gly Gly Phe Met Val Arg Gln Arg Ser Arg Thr Gly Gln Gly Lys 340 345
Thr Lys Arg Ser Val Ile Arg Lys Asp Ser Ser Gly Ser Ile Ser Glu 365 355
Gln Leu Pro Cys Arg Asp Asp Gly Trp Ser Glu Gln Leu Pro Asp Thr 370 375
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Lys Arg Tyr Arg Lys Arg Lys Asn Lys Leu Glu Glu Thr Phe P10 A10 415
Tyr Leu Gln Glu Ala Phe Phe Gly Lys Asp Leu Leu Asp Thr Ser Arg 430 420
Gln Ser Lys Ile Ser Leu Asp Asn Leu Ser Glu Asp Gly Ala Gln Leu 445 435
Leu Tyr Lys Thr Asn Met Asn Thr Gly Phe Leu Asp Pro Ser Leu Asp 460
Pro Leu Leu Ser Ser Ser Ala Pro Thr Lys Ser Gly Thr His Gly 480 475 465
Pro Ala Asp Asp Pro Leu Ala Asp Ile Ser Glu Val Leu Asn Thr Asp 495
Asp Asp Ile Leu Gly Ile Ile Ser Asp Asp Leu Ala Lys Ser Val Asp 500 500
His Ser Asp Ile Gly Pro Val Thr Asp Asp Pro Ser Ser Leu Pro Gln 525
Pro Asn Val Asn Gln Ser Ser Arg Pro Leu Ser Glu Glu Gln Leu Asp 540
Gly Ile Leu Ser Pro Glu Leu Asp Lys Met Val Thr Asp Gly Ala Ile 560 545
Leu Gly Lys Leu Tyr Lys Ile Pro Glu Leu Gly Gly Lys Asp Val Glu 575 565
Asp Leu Phe Thr Ala Val Leu Ser Pro Ala Asn Thr Gln Pro Thr Pro 580 580

Leu Pro Gln Pro Pro Pro Pro Thr Gln Leu Leu Pro Ile His Asn Gln 595 600 607
Asp Ala Phe Ser Arg Met Pro Leu Met Asn Gly Leu Ile Gly Ser Ser 610 615 610 617
Pro His Leu Pro His Asn Ser Leu Pro Pro Gly Ser Gly Leu Gly Thr 640 635 630 630 630 630 630 630 630
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Arg Ser Thr Leu Lys Trp Glu Lys Glu Glu Ala Leu Gly Glu Met Ala 690 695
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Glu Glu Phe Pro Asp Trp Thr Thr Arg Val Lys Gln Ile Ala Lys Leu 735 725
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Arg Ile Asp Ser Glu Leu Phe Lys Asp Pro Leu Lys Gln Arg Glu Sel 790 795
Glu His Glu Gln Glu Trp Lys Phe Arg Gln Gln Met Arg Gln Lys Ser 815
Lys Gln Gln Ala Lys Ile Glu Ala Thr Gln Lys Leu Glu Gln Val Lys 820 825
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Leu Val Gln Ser Gly Ser Asp Thr Pro Ser Ser Gly Ile Gln Ser Pro 850
Leu Thr Pro Gln Pro Gly Asn Gly Asn Met Ser Pro Ala Gln Ser Phe 875 865

His Lys Glu Leu Phe Thr Lys Gln Pro Pro Ser Thr Pro Thr Ser Thr 885 Ser Ser Asp Asp Val Phe Val Lys Pro Gln Ala Pro Pro Pro Pro Ala Pro Ser Arg Ile Pro Ile Gln Asp Ser Leu Ser Gln Ala Gln Thr Ser Gln Pro Pro Ser Pro Gln Val Phe Ser Pro Gly Ser Ser Asn Ser Arg Pro Pro Ser Pro Met Asp Pro Tyr Ala Lys Met Val Gly Thr Pro 950 Arg Pro Pro Pro Val Gly His Ser Phe Ser Arg Arg Asn Ser Ala Ala Pro Val Glu Asn Cys Thr Pro Leu Ser Ser Val Ser Arg Pro Leu Gln 980 Met Asn Glu Thr Thr Ala Asn Arg Pro Ser Pro Val Arg Asp Leu Cys Ser Ser Ser Thr Thr Asn Asn Asp Pro Tyr Ala Lys Pro Pro Asp Thr Pro Arg Pro Val Met Thr Asp Gln Phe Pro Lys Ser Leu Gly Leu Ser 1030 Arg Ser Pro Val Val Ser Glu Gln Thr Ala Lys Gly Pro Ile Ala Ala Gly Thr Ser Asp His Phe Thr Lys Pro Ser Pro Arg Ala Asp Val Phe 1060 Gln Arg Gln Arg Ile Pro Asp Ser Tyr Ala Arg Pro Leu Leu Thr Pro Ala Pro Leu Asp Ser Gly Pro Gly Pro Phe Lys Thr Pro Met Gln Pro 1095 Pro Pro Ser Ser Gln Asp Pro Tyr Gly Ser Val Ser Gln Ala Ser Arg 1090 1110 Arg Leu Ser Val Asp Pro Tyr Glu Arg Pro Ala Leu Thr Pro Arg Pro 1125 Ile Asp Asn Phe Ser His Asn Gln Ser Asn Asp Pro Tyr Ser Gln Pro 1140 Pro Leu Thr Pro His Pro Ala Val Asn Glu Ser Phe Ala His Pro Ser

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	Ser Gln Ser Ser Gly Thr Ala Arg Ser Asn Thr Asp Pro Tyr Ser Gln 1215 1205
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	Pro Gln Thr Pro Arg Pro Ser Thr Gln Thr Asp Leu Phe Val Thr Pro 1245 1235
	Val Thr Asn Gln Arg His Ser Asp Pro Tyr Ala His Pro Pro Gly Thr  1260 1250  The Gar Clar Pro Pro Ala Thr Pro
	Pro Arg Pro Gly Ile Ser Val Pro Tyr Ser Gln Pro Pro Ala Thr Pro 1280 1275 1265
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- Gly Arg Gln Glu Lys Gly Ser Gln Asp Ser Pro Ala Val Pro His Pro 1460 1460
- Gly Pro Leu Gln His Trp Gln Pro Glu Asn Val Asn Gln Ala Phe Thr 1485
- Arg Pro Pro Pro Pro Tyr Pro Gly Asn Ile Arg Ser Pro Val Ala Pro 1490
- Pro Leu Gly Pro Arg Tyr Ala Val Phe Pro Lys Asp Gln Arg Gly Pro 1520 1505
- Tyr Pro Pro Asp Val Ala Ser Met Gly Met Arg Pro His Gly Phe Arg 1535
- Phe Gly Phe Pro Gly Gly Ser His Gly Thr Met Pro Ser Gln Glu Arg 1540 1545
- Phe Leu Val Pro Pro Gln Gln Ile Gln Gly Ser Gly Val Ser Pro Gln 1555
- Leu Arg Arg Ser Val Ser Val Asp Met Pro Arg Pro Leu Asn Asn Ser 1570 1570 - Classer
- Gln Met Asn Asn Pro Val Gly Leu Pro Gln His Phe Ser Pro Gln Ser 1595 1590 1595
- Leu Pro Val Gln Gln His Asn Ile Leu Gly Gln Ala Tyr Ile Glu Leu 1615
- Arg His Arg Ala Pro Asp Gly Arg Gln Arg Leu Pro Phe Ser Ala Pro 1620
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- Ile Pro Arg Pro Asp Phe Pro Gly Pro Arg His Thr Asp Pro Met Arg 1650 1650
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- Ser Ser Met Val Met Arg Thr Leu Asn His Pro Leu Gly Gly Glu Phe 1700
- Ser Glu Ala Pro Leu Ser Thr Ser Val Pro Ser Glu Thr Thr Ser Asp 1715
- Asn Leu Gln Ile Thr Thr Gln Pro Ser Asp Gly Leu Glu Glu Lys Leu 1730

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Gly Val Glu Val Lys Asp Leu Asp Asp Glu Asp Leu Glu Asn Leu Asn 1775
Leu Asp Thr Glu Asp Gly Lys Val Val Glu Leu Asp Thr Leu Asp Asn 1780 1785
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Phe Asp Ile Ile Ala Tyr Thr Asp Pro Glu Leu Asp Met Gly Asp Lys 1810 1815
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Asp Asn Gln Cys Val Ser Val Glu Pro Lys Lys Glu Gln Glu Asn 1845 1850
Lys Thr Leu Val Leu Ser Asp Lys His Ser Pro Gln Lys Lys Ser Thr 1860 1865 1870
Val Thr Asn Glu Val Lys Thr Glu Val Leu Ser Pro Asn Ser Lys Val 1875 1880 1885
Glu Ser Lys Cys Glu Thr Glu Lys Asn Asp Glu Asn Lys Asp Asn Val 1890 1895 1900
Asp Thr Pro Cys Ser Gln Ala Ser Ala His Ser Asp Leu Asn Asp Gly 1905 1910 1915 1920
Glu Lys Thr Ser Leu His Pro Cys Asp Pro Asp Leu Phe Glu Lys Arg 1930 1935
Thr Asn Arg Glu Thr Ala Gly Pro Ser Ala Asn Val Ile Gln Ala Ser 1940 1945
Thr Gln Leu Pro Ala Gln Asp Val Ile Asn Ser Cys Gly Ile Thr Gly 1965
Ser Thr Pro Val Leu Ser Ser Leu Leu Ala Asn Glu Lys Ser Asp Asn 1970 1975 1980
Ser Asp Ile Arg Pro Ser Gly Ser Pro Pro Pro Pro Thr Leu Pro Ala 1985 1990 1995 2000
Ser Pro Ser Asn His Val Ser Ser Leu Pro Pro Phe Ile Ala Pro Pro 2015
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Arg Val Asn His Val Phe Ser Gln Gly Val Gln Val Asn Pro Gly Leu 2035
Ile Pro Gly Gln Ser Thr Val Asn His Ser Leu Gly Thr Gly Lys Pro 2050 2060
Ala Thr Gln Thr Gly Pro Gln Thr Ser Gln Ser Gly Thr Ser Ser Met 2080 2075
Ser Gly Pro Gln Gln Leu Met Ile Pro Gln Thr Leu Ala Gln Gln Asn 2095
Arg Glu Arg Pro Leu Leu Leu Glu Glu Gln Pro Leu Leu Gln Asp 2100 2100
Leu Leu Asp Gln Glu Arg Gln Glu Gln Gln Gln Arg Gln Met Gln 2125 2115
Ala Met Ile Arg Gln Arg Ser Glu Pro Phe Phe Pro Asn Ile Asp Phe 2130 2135 2140
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Gly Ile Asn Lys Val Met Ala Gln Asn Asn Leu Gly Met Pro Pro Met 2175 2165 2170 2175
Val Met Ser Arg Phe Pro Phe Met Gly Gln Val Val Thr Gly Thr Gln 2180 2185
Asn Ser Glu Gly Gln Asn Leu Gly Pro Gln Ala Ile Pro Gln Asp Gly 2195 2200 2205
Ser Ile Thr His Gln Ile Ser Arg Pro Asn Pro Pro Asn Phe Gly Pro 2210 2215 2220
Gly Phe Val Asn Asp Ser Gln Arg Lys Gln Tyr Glu Glu Trp Leu Gln 2240 2225
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Ile Gly Ala His Arg Lys Ser Lys Lys Ala Leu Ser Ala Lys Gln Arg 2260 2265 2270
Thr Ala Lys Lys Ala Gly Arg Glu Phe Pro Glu Glu Asp Ala Glu Gln 2285
Leu Lys His Val Thr Glu Gln Gln Ser Met Val Gln Lys Gln Leu Glu 2290 2295 2300
Gln Ile Arg Lys Gln Gln Lys Glu His Ala Glu Leu Ile Glu Asp Tyr 2305 2310 2315 2320

Arg Ile Lys Gln Gln Gln Cys Ala Met Ala Pro Pro Thr Met Met Pro Ser Val Gln Pro Gln Pro Pro Leu Ile Pro Gly Ala Thr Pro Pro Thr Met Ser Gln Pro Thr Phe Pro Met Val Pro Gln Gln Leu Gln His Gln Gln His Thr Thr Val Ile Ser Gly His Thr Ser Pro Val Arg Met Pro Ser Leu Pro Gly Trp Gln Pro Asn Ser Ala Pro Ala His Leu Pro Leu Asn Pro Pro Arg Ile Gln Pro Pro Ile Ala Gln Leu Pro Ile Lys Thr Cys Thr Pro Ala Pro Gly Thr Val Ser Asn Ala Asn Pro Gln Ser Gly Pro Pro Pro Arg Val Glu Phe Asp Asp Asn Asn Pro Phe Ser Glu Ser Phe Gln Glu Arg Glu Arg Lys Glu Arg Leu Arg Glu Gln Glu Arg Gln Arg Ile Gln Leu Met Gln Glu Val Asp Arg Gln Arg Ala Leu Gln Gln Arg Met Glu Met Glu Gln His Gly Met Val Gly Ser Glu Ile Ser Ser Ser Arg Thr Ser Val Ser Gln Ile Pro Phe Tyr Ser Ser Asp Leu Pro Cys Asp Phe Met Gln Pro Leu Gly Pro Leu Gln Gln Ser Pro Gln His Gln Gln Gln Met Gly Gln Val Leu Gln Gln Gln Asn Ile Gln Gln Gly Ser Ile Asn Ser Pro Ser Thr Gln Thr Phe Met Gln Thr Asn Glu Arg Arg Gln Val Gly Pro Pro Ser Phe Val Pro Asp Ser Pro Ser Ile Pro Val Gly Ser Pro Asn Phe Ser Ser Val Lys Gln Gly His Gly Asn Leu Ser Gly Thr Ser Phe Gln Gln Ser Pro Val Arg Pro Ser Phe

- Thr Pro Ala Leu Pro Ala Ala Pro Pro Val Ala Asn Ser Ser Leu Pro 2615 2610 Cys Gly Gln Asp Ser Thr Ile Thr His Gly His Ser Tyr Pro Gly Ser
- 2630 2625
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- Thr Tyr Ala Asn Ser Glu Val Asp Lys Leu Ser Met Glu Thr Pro Ala 2760
- Lys Thr Glu Glu Ile Lys Leu Glu Lys Ala Glu Thr Glu Ser Cys Pro 2775
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- Gly Asn Ala Val Ala Cys Pro Val Ser Ser Ala Gln Ser Pro Pro His 2810
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- Pro Pro Thr Pro Pro Ala Ser Leu Pro Pro Thr Pro Pro Pro Met Ala 3010 3015 3020
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- Lys Pro Phe Gln Leu Pro Phe Arg Pro Gln Asp Asp Leu Leu Ala Arg 3060 3065 3070
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- Thr Pro Pro His Asn Asn Gln Glu Glu Leu Arg Ile Gln Asp His Cys 3090 3095 3100
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- Lys Glu Glu Pro Pro Glu Pro Val Pro Ser Pro Ile Ile Pro Ile Leu 3140 3145 3150
- Pro Ser Thr Ala Gly Lys Ser Ser Glu Ser Arg Arg Asn Asp Ile Lys 3155 3160 3165
- Thr Glu Pro Gly Thr Leu Tyr Phe Ala Ser Pro Phe Gly Pro Ser Pro 3170 3175 3180

Asn Gly Pro Arg Ser Gly Leu Ile Ser Val Ala Ile Thr Leu His Pro 3190 3195 3185 Thr Ala Ala Glu Asn Ile Ser Ser Val Val Ala Ala Phe Ser Asp Leu 3210 3205 Leu His Val Arg Ile Pro Asn Ser Tyr Glu Val Ser Ser Ala Pro Asp 3225 Val Pro Ser Met Gly Leu Val Ser Ser His Arg Ile Asn Pro Gly Leu 3240 3235 Glu Tyr Arg Gln His Leu Leu Leu Arg Gly Pro Pro Pro Gly Ser Ala 3255 Asn Pro Pro Arg Leu Val Ser Ser Tyr Arg Leu Lys Gln Pro Asn Val 3275 3265 3270 Pro Phe Pro Pro Thr Ser Asn Gly Leu Ser Gly Tyr Lys Asp Ser Ser 3285 His Gly Ile Ala Glu Ser Ala Ala Leu Arg Pro Gln Trp Cys Cys His 3305 Cys Lys Val Val Ile Leu Gly Ser Gly Val Arg Lys Ser Phe Lys Asp 3320 Leu Thr Leu Leu Asn Lys Asp Ser Arg Glu Ser Thr Lys Arg Val Glu 3335 Lys Asp Ile Val Phe Cys Ser Asn Asn Cys Phe Ile Leu Tyr Ser Ser 3355 3350 3345 Thr Ala Gln Ala Lys Asn Ser Glu Asn Lys Glu Ser Ile Pro Ser Leu 3370 3365 Pro Gln Ser Pro Met Arg Glu Thr Pro Ser Lys Ala Phe His Gln Tyr 3385 Ser Asn Asn Ile Ser Thr Leu Asp Val His Cys Leu Pro Gln Leu Pro 3400 3395 Glu Lys Ala Ser Pro Pro Ala Ser Pro Pro Ile Ala Phe Pro Pro Ala 3415 3420 Phe Glu Ala Ala Gln Val Glu Ala Lys Pro Asp Glu Leu Lys Val Thr 3435 3425 Val Lys Leu Lys Pro Arg Leu Arg Ala Val His Gly Gly Phe Glu Asp 3450 3445

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- Asp Pro Val Pro Lys Asp Tyr Arg Lys Cys Cys Phe Cys His Glu Glu 3505 3510 3515 3520
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- Ile Lys Ala Gln Cys Met Phe Phe Lys Asp Lys Thr Met Leu Cys Pro 3605 3610 3615
- Met His Lys Pro Lys Gly Ile His Glu Gln Glu Leu Ser Tyr Phe Ala 3620 3625 3630
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- Phe His Ser Pro Lys Ala Leu Phe Pro Val Gly Tyr Glu Ala Ser Arg 3685 3690 3695
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Pro Tyr Ser Lys Gln Phe Val His Ser Lys Ser Ser Gln Tyr Arg Lys 3860 3865 3870

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Gln Gly Leu Gly Leu Tyr Ala Ala Arg Asp Ile Glu Lys His Thr Met 3890 3895 3900

Val Ile Glu Tyr Ile Gly Thr Ile Ile Arg Asn Glu Val Ala Asn Arg 3905 3910 3915 3920

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Arg Tyr Ile Asn His Ser Cys Ala Pro Asn Cys Val Ala Glu Val Val 3955 3960 3965

Thr Phe Glu Arg Gly His Lys Ile Ile Ile Ser Ser Ser Arg Arg Ile 3970 3975 3980

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Trp Met Asn

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Pro Glu Asn Gln Ala Leu Ala Arg Phe Tyr Cys Tyr Thr Glu Arg Thr

Ile Ala Lys Arg Leu Val Leu Arg Arg Asp Pro Ser Val Lys Arg Thr

Leu Cys Arg Gly Cys Ser Ser Leu Leu Val Pro Gly Leu Thr Cys Thr

His Arg Gln Arg Arg Cys Arg Gly Gln Arg Trp Thr Val Gln Thr Cys

Leu Thr Cys Gln Arg Ser Gln Arg Phe Leu Asn Asp Pro Gly His Leu 120

Leu Trp Gly Asp Arg Pro Glu Ala Gln Leu Gly Ser Gln Ala Asp Ser 135 130

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Leu	Arg	His 35	Glu	Phe	Asp	Ser	Glu 40	His	Val	Pro	Asp	Leu 45	Thr	Lys	Glu
Pro	Tyr 50	Val	Gln	Asp	Ile	His 55	Ser	Val	Gly	Ser	Leu 60	Cys	Lys	Leu	Tyr
Phe 65	Arg	Glu	Leu	Pro	Asn 70	Pro	Leu	Leu	Thr	Tyr 75	Gln	Leu	Tyr	Glu	Lys 80
Phe	Ser	Asp	Ala	Val 85	Ser	Ala	Ala	Thr	Asp 90	Glu	Glu	Arg	Leu	Ile 95	Lys
Ile	His	Asp	Val 100	Ile	Gln	Gln	Leu	Pro 105	Pro	Pro	His	Tyr	Arg 110	Thr	Leu
Glu	Phe	Leu 115	Met	Arg	His	Leu	Ser 120	Leu	Leu	Ala	Asp	Tyr 125	Cys	Ser	Ile
Thr	Asn 130	Met	His	Ala	Lys	Asn 135	Leu	Ala	Ile	Val	Trp 140	Ala	Pro	Asn	Leu
Leu 145	Arg	Ser	Lys	Gln	Ile 150	Glu	Ser	Ala	Cys	Phe 155	Ser	Gly	Thr	Ala	Ala 160
Phe	Met	Glu	Val	Arg 165	Ile	Gln	Ser	Val	Val 170	Val	Glu	Phe	Ile	Leu 175	Asn
His	Val	Asp	Val 180	Leu	Phe	Ser	Gly	Arg 185	Ile	Ser	Met	Ala	Met 190	Gln	Glu
Gly	Ala	Ala 195	Ser	Leu	Ser	Arg	Pro 200	Lys	Ser	Leu	Leu	Val 205	Ser	Ser	Pro
Ser	Thr 210	Lys	Leu	Leu	Thr	Leu 215	Glu	Glu	Ala	Gln	Ala 220	Arg	Thr	Gln	Ala
Gln 225	Val	Asn	Ser	Pro	Ile 230	Val	Thr	Glu	Asn	Lys 235	Tyr	Ile	Glu	Val	Gly 240
Glu	Gly	Pro	Ala	Ala 245	Leu	Gln	Gly	Lys	Phe 250	His	Thr	Ile	Ile	Glu 255	Phe
Pro	Leu	Glu	Arg 260	Lys	Arg	Pro	Gln	Asn 265	Lys	Met	Lys	Lys	Ser 270	Pro	Val
Gly	Ser	Trp 275	Arg	Ser	Phe	Phe	Asn 280	Leu	Gly	Lys	Ser	Ser 285	Ser	Val	Ser
Lys	Arg	Lys	Leu	Gln	Arg	Asn	Glu	Ser	Glu	Pro	Ser	Glu	Met	Lys	Ala

	290	)				295	5				300	)			
Met 305	Ala	Leu	Lys	Gly	Gly 310		g Ala	Glu	Gly	7 Thr 315		ı Arç	g Ser	: Ala	Lys 320
Ser	Glu	Glu	. Ser	Leu 325		Ser	Leu	His	330		. Asp	Gly	/ Asp	Ser 335	_
Leu	. Phe	Arg	Pro 340	Arg	Arg	Pro	Arg	Ser 345		Ser	Asp	Ala	Leu 350		Ala
Ser	Phe	Asn 355	Gly	Glu	Met	Leu	Gly 360		Arg	Cys	Asn	Ser 365		Asp	Asn
Leu	Pro 370	His	Asp	Asn	Glu	Ser 375	Glu	Glu	Glu	Gly	Gly 380		Leu	His	Ile
Pro 385	Ala	Leu	Met	Ser	Pro 390		Ser	Ala	Glu	Asp 395		Asp	Leu	Ser	Pro 400
Pro	Asp	Ile	Gly	Val 405	Ala	Ser	Leu	Asp	Phe 410		Pro	Met	Ser	Phe 415	
Cys	Ser	Pro	Pro 420	Lys	Ala	Glu	Ser	Glu 425	Cys	Leu	Glu	Ser	Gly 430	Ala	Ser
Phe	Leu	Asp 435	Ser	Pro	Gly	Tyr	Ser 440	Lys	Asp	Lys	Pro	Ser 445	Ala	Asn	Lys
Lys	Asp 450	Ala	Glu	Thr	Gly	Ser 455	Ser	Gln	Cys	Gln	Thr 460	Pro	Gly	Ser	Thr
Ala 465	Ser	Ser	Glu	Pro	Val 470	Ser	Pro	Leu	Gln	Glu 475	Lys	Leu	Ser	Pro	Phe 480
Phe	Thr	Leu	Asp	Leu 485	Ser	Pro	Thr	Glu	Asp 490	Lys	Ser	Ser	Lys	Pro 495	Ser
Ser	Phe	Thr	Glu 500	Lys	Val	Val	Tyr	Ala 505	Phe	Ser	Pro	Lys	Ile 510	Gly	Arg
Lys	Leu	Ser 515	Lys	Ser	Pro	Ser	Met 520	Ser	Ile	Ser	Glu	Pro 525	Ile	Ser	Val
Thr	Leu 530	Pro	Pro	Arg	Val	Ser 535	Glu	Val	Ile	Gly	Thr 540	Val	Ser	Asn	Thr
Thr 545	Ala	Gln	Asn	Ala	Ser 550	Ser	Ser	Thr	Trp	Asp 555	Lys	Cys	Val	Glu	Glu 560
Arg	Asp	Ala	Thr	Asn 565	Arg	Ser	Pro	Thr	Gln 570	Ile	Val	Lys	Met	Lys 575	Thr
Asn	Glu	Thr	Val	Ala	Gln	Glu	Ala	Tyr	Glu	Ser	Glu	Val	Gln	Pro	Leu

			580					585					590		
Asp	Gln	Val 595	Ala	Ala	Glu	Glu	Val 600	Glu	Leu	Pro	Gly	Lys 605	Glu	Asp	Glr
Ser	Val 610	Ser	Ser	Ser	Gln	Ser 615	Lys	Ala	Val	Ala	Ser 620	Gly	Gln	Thr	Glr
Thr 625	Gly	Ala	Val	Thr	His 630	Asp	Pro	Pro	Gln	Asp 635	Ser	Val	Pro	Val	Ser 640
Ser	Val	Ser	Leu	Ile 645	Pro	Pro	Pro	Pro	Pro 650	Pro	Lys	Asn	Val	Ala 655	Arg
Met	Leu	Ala	Leu 660	Ala	Leu	Ala	Glu	Ser 665	Ala	Gln	Gln	Ala	Ser 670	Thr	Gln
Ser	Leu	Lys 675	Arg	Pro	Gly	Thr	Ser 680	Gln	Ala	Gly	Tyr	Thr 685	Asn	Tyr	Gly
Asp	Ile 690	Ala	Val	Ala	Thr	Thr 695	Glu	Asp	Asn	Leu	Ser 700	Ser	Ser	Tyr	Ser
Ala 705	Val	Ala	Leu	Asp	Lys 710	Ala	Tyr	Phe	Gln	Thr 715	Asp	Arg	Pro	Ala	Glu 720
Gln	Phe	His	Leu	Gln 725	Asn	Asn	Ala	Pro	Gly 730	Asn	Cys	Asp	His	Pro 735	Leu
Pro	Glu	Thr	Thr 740	Ala	Thr	Gly	Asp	Pro 745	Thr	His	Ser	Asn	Thr 750	Thr	Glu
Ser	Gly	Glu 755	Gln	His	His	Gln	Val 760	Asp	Leu	Thr	Gly	Asn 765	Gln	Pro	His
Gln	Ala 770	Tyr	Leu	Ser	Gly	Asp 775	Pro	Glu	Lys	Ala	Arg 780	Ile	Thr	Ser	Val
Pro 785	Leu	Asp	Ser	Glu	Lys 790	Ser	Asp	Asp	His	Val 795	Ser	Phe	Pro	Glu	Asp 800
Gln	Ser	Gly	Lys	Asn 805	Ser	Met	Pro	Thr	Val 810	Ser	Phe	Leu	Asp	Gln 815	Asp
Gln	Ser	Pro	Pro 820	Arg	Phe	Tyr	Ser	Gly 825	Asp	Gln	Pro	Pro	Ser 830	Tyr	Leu
Gly	Ala	Ser 835	Val	Asp	Lys	Leu	His 840	His	Pro	Leu	Glu	Phe 845	Ala	Asp	Lys
Ser	Pro 850	Thr	Pro	Pro	Asn	Leu 855	Pro	Ser	Asp	Lys	Ile 860	Tyr	Pro	Pro	Ser
Gly	Ser	Pro	Glu	Glu	Asn	Thr	Ser	Thr	Ala	Thr	Met	Thr	Tvr	Met	Thr

865					870					875					880
Thr	Thr	Pro	Ala	Thr 885		Gln	Met	Ser	Thr 890		Glu	Ala	Ser	Trp 895	Asp
Val	Ala	Glu	Gln 900		Thr	Thr	Ala	Asp 905		Ala	Ala	Ala	Thr 910		Gln
Arg	Thr	His 915		Thr	Asn	Arg	Pro 920		Pro	Pro	Pro	Pro 925		Gln	Arg
Ser	Ala 930	Glu	Gln	Pro	Pro	Val 935	Val	Gly	Gln	Val	Gln 940	Ala	Ala	Thr	Asn
Ile 945		Leu	Asn	Asn	Ser 950	His	Lys	Val	Gln	Gly 955		Val	Pro	Val	Pro 960
Glu	Arg	Pro	Pro	Glu 965		Arg	Ala	Met	Asp 970	Asp	Pro	Ala	Ser	Ala 975	Phe
Ile	Ser	Asp	Ser 980	Gly	Ala	Ala	Ala	Ala 985	Gln	Cys	Pro	Met	Ala 990	Thr	Ala
Val	Gln	Pro 995	Gly	Leu	Pro	Glu	Lys 100		Arg	Asp	Gly	Ala 100		Val	Pro
Leu	Leu 1010		Leu	Arg	Ala	Glu 101		Val	Pro	Ala	His 102		Cys	Gly	Phe
Pro 102		Pro	Leu	Pro	Pro 1030		Arg	Met	Met	Glu 103		Lys	Met	Ile	Ala 1040
Ala	Ile	His	Ser	Ser 104		Ala	Asp	Ala	Thr 1050		Ser	Ser	Asn	Tyr 105	His 5
Ser	Phe	Val	Thr 1060		Ser	Ser	Thr	Ser 106		Asp	Asp	Ala	Leu 1070		Leu
Pro	Leu	Pro 1075	Val 5	Pro	Gln	Pro	Lys 1080		Ala	Ser	Gln	Lys 108		Val	Tyr
Ser	Ser 1090		Ala	Arg	Pro	Asp 1095		Thr	Thr	Glu	Pro 1100		Gly	Pro	Asp
Asn 1105		Leu	His	Phe	Asn 1110		Thr	Pro	Asn	Cys 1115		Tyr	Arg	Pro	Gln 1120
Ser	Val	Pro	Pro	His 1125	His	Asn	Lys	Leu	Glu 1130		His	Gln	Val	Tyr 1135	-
Ala	Arg	Ser	Glu 1140		Pro	Ala	Ser	Met 1145		Leu	Arg	Tyr	Asn 1150		Tyr
Val	Ala	Pro	Gly	Arg	Asn	Ala	Ser	Gly	His	His	Ser	Lys	Pro	Cys	Ser

		115	55				116	0				116	55		
Arg	Val 117		ı Tyr	Val	. Ser	Ser 117		Ser	Ser	Ser	Val 118	-	Asn	Thr	Cys
Tyr 118	Pro 5	Glu	ı Asp	Ile	Pro 119		Tyr	Pro	Thr	11e		Arg	Val	Gln	Ser 1200
Leu	His	Ala	Pro	Pro 120	Ser 5	Ser	Met	Ile	Arg 121		Val	Pro	Ile	Ser 121	-
Thr	Glu	Val	Pro 122		Asp	Asp	Glu	Pro 122		Tyr	Cys	Pro	Arg 123		Leu
Tyr	Gln	Tyr 123	Lys 5	Pro	Tyr	Gln	Ser 124		Gln	Ala	Arg	Ser 124		Tyr	His
Val	Thr 125	Gln O	Leu	Gln	Pro	Tyr 125		Glu	Asn	Gly	Arg 126		His	Tyr	Arg
Tyr 126		Pro	Tyr	Ser	Ser 1270		Ser	Ser	Ser	Tyr 127		Ser	Pro	Asp	Gly 1280
Ala	Leu	Cys	Asp	Val 128	Asp 5	Ala	Tyr	Gly	Thr 129		Gln	Leu	Arg	Pro 129	
His	Arg	Leu	Pro 1300		Arg	Asp	Phe	Ala 130		Tyr	Asn	Pro	Arg 131		Gln
Gly	Lys	Ser 131	Leu 5	Tyr	Ser	Tyr	Ala 1320		Leu	Ala	Pro	Arg 132		Arg	Ala
Asn	Val 1330	Thr	Gly	Tyr	Phe	Ser 1335		Asn	Asp	His	Asn 1340		Val	Ser	Met
Pro 1345	Pro	Ala	Ala	Asp	Val 1350		His	Thr	Tyr	Thr 1355		Trp	Asp	Leu	Glu 1360
Asp	Met	Glu	Lys	Tyr 1365	Arg	Met	Gln	Ser	Ile 1370		Arg	Glu	Ser	Arg 1375	
Arg	Gln	Lys	Val 1380		Gly	Pro	Val	Met 1385		Gln	Tyr	Asp	Asn 1390		Thr
Pro	Ala	Val 1395	Gln	Asp	Asp	Leu	Gly 1400		Ile	Tyr	Val	Ile 1405		Leu	Arg
Ser	Lys 1410		Asp	Pro	Gly	Lys 1415		Gly	Leu	Leu	Ser 1420		Ala	Glu	Gly
Lys 1425	Glu	Ser	Arg	His	Ala 1430		Lys	Ala	Ile	Ser 1435		Glu	Gly		Asp 1440
Ara	Phe	Tvr	Ara	Ara	His	Pro	Gl 11	<b>Δ</b> 1 =	Gl 11	Mot	Λen	7 ~~	חות	ui a	II

1445 1450 1455 His Gly Gly His Gly Ser Thr Gln Pro Glu Lys Pro Ser Leu Pro Gln 1460 1465 Lys Gln Ser Ser Leu Arg Ser Arg Lys Leu Pro Asp Met Gly Cys Ser Leu Pro Glu His Arg Ala His Gln Glu Ala Ser His Arg Gln Phe Cys 1490 1495 Glu Ser Lys Asn Gly Pro Pro Tyr Pro Gln Gly Ala Gly Gln Leu Asp 1510 Tyr Gly Ser Lys Gly Ile Pro Asp Thr Ser Glu Pro Val Ser Tyr His 1530 Asn Ser Gly Val Lys Tyr Ala Ala Ser Gly Gln Glu Ser Leu Arg Leu 1540 1545 1550 Asn His Lys Glu Val Arg Leu Ser Lys Glu Met Glu Arg Pro Trp Val 1560 Arg Gln Pro Ser Ala Pro Glu Lys His Ser Arg Asp Cys Tyr Lys Glu 1570 1575 1580 Glu Glu His Leu Thr Gln Ser Ile Val Pro Pro Pro Lys Pro Glu Arg 1590 1595 Ser His Ser Leu Lys Leu His His Thr Gln Asn Val Glu Arg Asp Pro 1605 1610 Ser Val Leu Tyr Gln Tyr Gln Pro His Gly Lys Arg Gln Ser Ser Val 1620 1625 Thr Val Val Ser Gln Tyr Asp Asn Leu Glu Asp Tyr His Ser Leu Pro 1640 Gln His Gln Arg Gly Val Phe Gly Gly Gly Met Gly Thr Tyr Val 1650 1655 1660 Pro Pro Gly Phe Pro His Pro Gln Ser Arg Thr Tyr Ala Thr Ala Leu 1670 1675 Gly Gln Gly Ala Phe Leu Pro Ala Glu Leu Ser Leu Gln His Pro Glu 1685 1690 Thr Gln Ile His Ala Glu

1700

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<400> 435

Pro Phe Gln Gln Val Gly Arg Cys Asn Pro Ser Pro Gln Thr Arg Pro 5 10 15

Gly Pro Ala Ser Lys Val Lys Gln Asp Met Pro Pro Pro Gly Gly Tyr
20 25 30

Gly Pro Ile Asp Tyr Lys Arg Asn Leu Pro Arg Arg Gly Leu Ser Gly 35 40 45

Tyr Ser Met Leu Ala Ile Gly Ile Gly Thr Leu Ile Tyr Gly His Trp 50 55 60

Ser Ile Met Lys Trp Asn Arg Glu Arg Arg Arg Leu Gln Ile Glu Asp 65 70 75 80

Phe Glu Ala Arg Ile Ala Leu Leu Pro Leu Leu Gln Ala Glu Thr Asp 85 90 95

Arg Arg Thr Leu Gln Met Leu Arg Glu Asn Leu Glu Glu Glu Ala Ile 100 105 110

Ile Met Lys Asp Val Pro Asp Trp Lys Val Gly Glu Ser Val Phe His
115 120 125

Thr Thr Arg Trp Val Pro Pro Leu Ile Gly Glu Leu Tyr Gly Leu Arg 130 135 140

Thr Thr Glu Glu Ala Leu His Ala Ser His Gly Phe Met Trp Tyr Thr 145 150 155 160

<210> 436

<211> 396

<212> PRT

<213> Homo sapiens

<400> 436

Arg Ala Gln Glu Ala Ala Ala Ala Ala Ala Asp Gly Pro Pro Ala Ala 5 10 15

Asp Gly Glu Asp Gly Gln Asp Pro His Ser Lys His Leu Tyr Thr Ala 20 25 30

Asp Met Phe Thr His Gly Ile Gln Ser Ala Ala His Phe Val Met Phe 35 40 45

Phe Ala Pro Trp Cys Gly His Cys Gln Arg Leu Gln Pro Thr Trp Asn 50 55 60

Asp Leu Gly Asp Lys Tyr Asn Ser Met Glu Asp Ala Lys Val Tyr Val 65 70 75 80

Ala	Lys	Val	Asp	Cys 85	Thr	Ala	His	Ser	Asp 90	Val	Cys	Ser	Ala	Gln 95	Gly
Val	Arg	Gly	Tyr 100	Pro	Thr	Leu	Lys	Leu 105	Phe	Lys	Pro	Gly	Gln 110	Glu	Ala
Val	Lys	Tyr 115	Gln	Gly	Pro	Arg	Asp 120	Phe	Gln	Thr	Leu	Glu 125	Asn	Trp	Met
Leu	Gln 130	Thr	Leu	Asn	Glu	Glu 135	Pro	Val	Thr	Pro	Glu 140	Pro	Glu	Val	Glu
Pro 145	Pro	Ser	Ala	Pro	Glu 150	Leu	Lys	Gln	Gly	Leu 155	Tyr	Glu	Leu	Ser	Ala 160
Ser	Asn	Phe	Glu	Leu 165	His	Val	Ala	Gln	Gly 170	Asp	His	Phe	Ile	Lys 175	Phe
Phe	Ala	Pro	Trp 180	Cys	Gly	His	Cys	Lys 185	Ala	Leu	Ala	Pro	Thr 190	Trp	Glu
Gln	Leu	Ala 195	Leu	Gly	Leu	Glu	His 200	Ser	Glu	Thr	Val	Lys 205	Ile	Gly	Lys
Val	Asp 210	Cys	Thr	Gln	His	Tyr 215	Glu	Leu	Cys	Ser	Gly 220	Asn	Gln	Val	Arg
Gly 225	Tyr	Pro	Thr	Leu	Leu 230	Trp	Phe	Arg	Asp	Gly 235	Lys	Lys	Val	Asp	Gln 240
Tyr	Lys	Gly	Lys	Arg 245	Asp	Leu	Glu	Ser	Leu 250	Arg	Glu	Tyr	Val	Glu 255	Ser
Gln	Leu	Gln	Arg 260	Thr	Glu	Thr	Gly	Ala 265	Thr	Glu	Thr	Val	Thr 270	Pro	Ser
Glu	Ala	Pro 275	Val	Leu	Ala	Ala	Glu 280	Pro	Glu	Ala	Asp	Lys 285	Gly	Thr	Val
Leu	Ala 290	Leu	Thr	Glu	Asn	Thr 295	Phe	Asp	Asp	Thr	Ile 300	Ala	Glu	Gly	Ile
Thr 305	Phe	Ile	Lys	Phe	Tyr 310	Ala	Pro	Trp	Cys	Gly 315	His	Cys	Lys	Thr	Leu 320
Ala	Pro	Thr	Trp	Glu 325	Glu	Leu	Ser	Lys	Lys 330	Glu	Phe	Pro	Gly	Leu 335	Ala
Gly	Val	Lys	Ile 340	Ala	Glu	Val	Asp	Cys 345	Thr	Ala	Glu	Arg	Asn 350	Ile	Cys
Ser	Lys	Tyr 355	Ser	Val	Arg	Gly	Tyr 360	Pro	Thr	Leu	Leu	Leu 365	Phe	Arg	Gly

Gly Lys Lys Val Ser Glu His Ser Gly Gly Arg Asp Leu Asp Ser Leu 370 380

His Arg Phe Val Leu Ser Gln Ala Lys Asp Glu Leu 385 390 395

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<211> 92

<212> PRT

<213> Homo sapiens

<400> 437

Ala Glu Met Asp Pro Leu Arg Ala Gln Gln Leu Ala Ala Glu Leu Glu 5 10 15

Val Glu Met Met Ala Asp Met Tyr Asn Arg Met Thr Ser Ala Cys His
20 25 30

Arg Lys Cys Val Pro Pro His Tyr Lys Glu Ala Glu Leu Ser Lys Gly
35 40 45

Glu Ser Val Cys Leu Asp Arg Cys Val Ser Lys Tyr Leu Asp Ile His 50 55 60

Glu Arg Met Gly Lys Lys Leu Thr Glu Leu Ser Met Gln Asp Glu Glu 65 70 75 80

Leu Met Lys Arg Val Gln Gln Ser Ser Gly Pro Ala 85 90

<210> 438

<211> 303

<212> PRT

<213> Homo sapiens

<400> 438

Lys Asn Pro Ala Lys Met Ser Leu Tyr Pro Ser Leu Glu Asp Leu Lys
5 10 15

Val Asp Lys Val Ile Gln Ala Gln Thr Ala Phe Ser Ala Asn Pro Ala 20 25 30

Asn Pro Ala Ile Leu Ser Glu Ala Ser Ala Pro Ile Pro His Asp Gly 35 40 45

Asn Leu Tyr Pro Arg Leu Tyr Pro Glu Leu Ser Gln Tyr Met Gly Leu 50 55 60

Ser Leu Asn Glu Glu Glu Ile Arg Ala Asn Val Ala Val Val Ser Gly
65 70 75 80

Ala Pro Leu Gln Gly Gln Leu Val Ala Arg Pro Ser Ser Ile Asn Tyr 85 Met Val Ala Pro Val Thr Gly Asn Asp Val Gly Ile Arg Arg Ala Glu 105 Ile Lys Gln Gly Ile Arg Glu Val Ile Leu Cys Lys Asp Gln Asp Gly Lys Ile Gly Leu Arg Leu Lys Ser Ile Asp Asn Gly Ile Phe Val Gln 135 Leu Val Gln Ala Asn Ser Pro Ala Ser Leu Val Gly Leu Arg Phe Gly 150 155 Asp Gln Val Leu Gln Ile Asn Gly Glu Asn Cys Ala Gly Trp Ser Ser 165 170 Asp Lys Ala His Lys Val Leu Lys Gln Ala Phe Gly Glu Lys Ile Thr 185 Met Thr Ile Arg Asp Arg Pro Phe Glu Arg Thr Ile Thr Met His Lys 200 Asp Ser Thr Gly His Val Gly Phe Ile Phe Lys Asn Gly Lys Ile Thr Ser Ile Val Lys Asp Ser Ser Ala Ala Arg Asn Gly Leu Leu Thr Glu 230 235 His Asn Ile Cys Glu Ile Asn Gly Gln Asn Val Ile Gly Leu Lys Asp 245 250 255 Ser Gln Ile Ala Asp Ile Leu Ser Thr Ser Gly Thr Val Val Thr Ile 260 Thr Ile Met Pro Ala Phe Ile Phe Glu His Ile Ile Lys Arg Met Ala 280 Pro Ser Ile Met Lys Ser Leu Met Asp His Thr Ile Pro Glu Val 295 300 <210> 439 <211> 378 <212> PRT <213> Homo sapiens <400> 439 Val Val Pro Ser Thr Lys Asp Phe Leu Val Gly Val Lys Gly Ser Gly

Gly His Arg Gly Gly Glu Met Ala Phe Ser Gly Ser Gln Ala Pro

5

Tyr Leu Ser Pro Ala Val Pro Phe Ser Gly Thr Ile Gln Gly Gly Leu 35 40 45

Gln Asp Gly Leu Gln Ile Thr Val Asn Gly Thr Val Leu Ser Ser Ser

Gly Thr Arg Phe Ala Val Asn Phe Gln Thr Gly Phe Ser Gly Asn Asp
65 70 75 80

Ile Ala Phe His Phe Asn Pro Arg Phe Glu Asp Gly Gly Tyr Val Val
85 90 95

Cys Asn Thr Arg Gln Asn Gly Ser Trp Gly Pro Glu Glu Arg Lys Thr
100 105 110

His Met Pro Phe Gln Lys Gly Met Pro Phe Asp Leu Cys Phe Leu Val 115 120 125

Gln Ser Ser Asp Phe Lys Val Met Val Asn Gly Ile Leu Phe Val Gln 130 135 140

Tyr Phe His Arg Val Pro Phe His Arg Val Asp Thr Ile Ser Val Asn 145 150 155 160

Gly Ser Val Gln Leu Ser Tyr Ile Ser Phe Gln Asn Pro Arg Thr Val 165 170 175

Pro Val Gln Pro Ala Phe Ser Thr Val Pro Phe Ser Gln Pro Val Cys 180 185 190

Phe Pro Pro Arg Pro Arg Gly Arg Arg Gln Lys Pro Pro Gly Val Trp 195 200 205

Pro Ala Asn Pro Ala Pro Ile Thr Gln Thr Val Ile His Thr Val Gln 210 215 220

Ser Ala Pro Gly Gln Met Phe Ser Thr Pro Ala Ile Pro Pro Met Met 225 230 235 240

Tyr Pro His Pro Ala Tyr Pro Met Pro Phe Ile Thr Thr Ile Leu Gly 245 250 255

Gly Leu Tyr Pro Ser Lys Ser Ile Leu Leu Ser Gly Thr Val Leu Pro 260 265 270

Ser Ala Gln Arg Phe His Ile Asn Leu Cys Ser Gly Asn His Ile Ala 275 280 285

Phe His Leu Asn Pro Arg Phe Asp Glu Asn Ala Val Val Arg Asn Thr 290 295 300

Gln Ile Asp Asn Ser Trp Gly Ser Glu Glu Arg Ser Leu Pro Arg Lys 305 310 315 320

Met Pro Phe Val Arg Gly Gln Ser Phe Ser Val Trp Ile Leu Cys Glu 325 330 335

Ala His Cys Leu Lys Val Ala Val Asp Gly Gln His Leu Phe Glu Tyr 340 345 350

Tyr His Arg Leu Arg Asn Leu Pro Thr Ile Asn Arg Leu Glu Val Gly 355 360 365

Gly Asp Ile Gln Leu Thr His Val Gln Thr 370 375

<210> 440 <211> 2239 <212> DNA

<213> Homo sapiens

<400> 440

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tacctattgg ggttaggaga aaagactaga caattactat gtggtcattc tctacaacat 2040 atgttagcac ggcaaagaac cttcaaattg aagactgaga tttttctgta tatatgggtt 2100 ttgtaaagat ggtttacac actacagatg tctatactgt gaaaagtgtt ttcaattctg 2160 aaaaaaaagca tacatcatga ttatggcaaa gaggagagaa ggtagagctg ttcttaaatt 2220 tattaaaaaa aaaaaaaaa 2239

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  <213> Homo sapiens
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<223> n = A, T, C or G
<400> 453
gaattcgaac cccttcggaa ggccaagggn ntagaaggng gctccggccc cagctgtcgt 60
gaagaagcag gaggctaaga aagtggtgaa tcccctgttt gagaaaaggc ctaagaattt 120
tggcattgga caggacatcc agcccaaaag agacctcacc cgctttgtga aatggccccg 180
ctatatcagg ttgcagcggc agagagccat cctctataag cggctgaaag tgcctcctgc 240
gattaaccag ttcacccagg ccctggaccg ccaaacagct actcagctgc ttaagctggc 300
ccacaagtac agaccagaga caaagcaaga gaagaagcag agactgttgg cccgggccga 360
gaagaagget getggeaaag gggaegteee aacgaagaga ceaectgtee ttegageagg 420
agttaacacc cgtcaccacc ttggtggaga acaagaaagc tcagctggtg gtgattgcac 480
acgacgtgga teceategag etggttgtet tettgeetge eetgtgtegt aaaatggggg 540
tcccttactg cattatcaag ggaaaggcaa gactgggacg tctagtccac aggaagacct 600
gcaccactgt cgccttccac aggtgaactc
                                                                   630
```

```
<211> 677
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(677)
<223> n = A, T, C or G
<400> 454
gaattcgaac cccttcgccc gcatgcggna catccccttg gccccagggt cagactggcg 60
cgatctgccc aacatcgagg tgcggctctc agacggcacc atggccagga agctgcggta 120
tacccaccat gacaggaaga acggccgcag cagctctggg gccctccgtg gggtctgctc 180
ctgcgtggaa gccggcaaag cctgcgaccc cgcagccagg cagttcaaca ccctcatccc 240
etggtgeetg ceceacaceg ggaaceggea caaceactgg getggeetet atggaagget 300
cgagtgggac ggcttcttca gcacaaccgt caccaacccc gagcccatgg gcaagcaggg 360
eegegtgete cacecagage ageacegtgt ggtgagegtg egggagtgtg eeegeteeea 420
gggcttccct gacacctacc ggctcttcgg caacatcctg gacaagcacc ggcaggtggg 480
caatgccgtg ccaccgcccc tggcaaagcc attggcttgg agatcaagct ttgtattgtt 540
ggccaaagcc cgagagagtg cctcagctaa aataaaggag gaggaagctg ctaaggacta 600
gttctgcctt cccgtcaccc ctgtttctgg caccaggaat cccccacaat gcacttgatg 660
gtggggtttt aacatgt
                                                                   677
<210> 455
<211> 598
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(598)
<223> n = A, T, C or G
<400> 455
ttttttggtt tataggagag atttatttga agaaatatta caacatataa aaactacata 60
aagtettaat tteeacteat aeagtggtag atttgatata atgeataata aaaaaetttt 120
aaaatccaga atgcacaaag tactgcacaa tttgatcact aaatcattag ttgataagcg 180
aacctcacac aacagcttca tgtcagccaa ggccacaaac accatgtacc acacatgtga 240
acggacagat tgacatgtta aaaacacaac atcagtgcat gttggggatt cctggtgcca 300
gaaacagggg tgacgggagg gcagaactag tccttagcag cttcctcctc ctttatttta 360
gctgaggcac tctctcgggc tttggccaac atacaaagct tgatctccaa gccaatggct 420
ttggccaggg gcggtggcac ggcattgccc acctgccggt gcttngtcca ggatgttgcc 480
cgaagagccg gtaggtggtc aagggaagcc cctggggaag cgggcacact cccggacgct 540
naccacacgg tgctgntttt gggtggagca ccgcggcctt gcttgcccat gggctcgg
<210> 456
<211> 574
<212> DNA
<213> Homo sapiens
<400> 456
ggaattcgaa ccccttcggg gcggggagcc ccgtagaacc gagggggtcg gcccgggggt 60
cccgggggag gtggagatgg tgaaggggca gccgttcgac gtgggcccgc gctacacgca 120
gttgcagtac atcggcgagg gcgcgtacgg catggtcagc tcggcctatg accacgtgcg 180
```

```
caagactcgc gtggccatca agaagatcag ccccttcgaa catcagacct actgccagcg 240
cacgeteegg gagateeaga teetgetgeg etteegeeat gagaatgtea teggeateeg 300
agacattetg egggegteea ceetggaage catgagagat gtetacattg tgeaggaeet 360
gatggagact gacctgtaca agttgctgaa aagccagcag ctgagcaatg accatatctg 420
ctactteete taccagatee tgeggggeet caagtacate caeteegeea aegtgeteea 480
ccgagatcta aagccctcca acctgcttca tcaacaccac ctggcgacct ttaaaatttg 540
                                                                 574
tgaatttccg gcctggcccc cggattgccc gaat
<210> 457
<211> 546
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(546)
<223> n = A, T, C or G
<400> 457
ttttttgaca catctctata tttatatatt agacgggtca gggaggtggc aggggcgccg 60
ggctctccac gccccccagc tccacttctg ctcaccacac acagaagcag cgagggcacg 120
cccttgagac taaggaatgt tccttcaggg aaactagggt ggggtttgaa tganatgagg 240
ggggcaggca tggccctgag tccctactca gcgcccccca ccctccacct ctgcccttca 300
gcaggttggg gcagccagaa cccttccatt ccagaactgc cagagactgg gacgctgggg 360
aaggtaaggg cgcagcagca gcagcgggag attgaactgg ggccacctga gctcccgagg 420
ccccgtgggg agggcgggtg gggaggaaaa ggccttggcc tgcctgaagc tggaggcctc 480
agcaaaggag agaggtggcc aggcccatgc tccaccccgg cctgggctgc caanggtccc 540
gggctg
<210> 458
<211> 674
<212> DNA
<213> Homo sapiens
<400> 458
gaattcgaac cccttcggta ttattaagaa ctaagagaat agcttgccag atacaaatgg 60
aaacacette caaatgagte ggagaaaatg tettgeagta ttatgggtaa aatageaaag 120
agcttgggaa tacagtttgc taatatcaag tccttaacaa cgaccattct tcattcaaga 180
ttagttgtgt ataaatacat gcttcttcag gagttgactt agaaaacaag caaacaaaca 240
aacatcagaa actatttaca actgggagca atccttgaag aacataaaga atataaatat 300
caacaaaggc tgaaaactct tttttagatt aaagatcaaa tggacatgtc atcggaatgt 360
attgtatggc tcttgattaa atcctggagc aaagtggaga gtgaggaaca actgtaaaga 420
atqtqaatac qqactqtqta ttaqataaca qtaccataaa tttcctqqat qqqataatta 480
tgttgtgact atgtaagaga atattttgcc cttagaagat atatgatgaa gcatttagaa 540
gtaaagtatc atgacatctt gcaaataact ttcaagtgat tcagccagat atataaaaat 600
tatatataac acattatata atttatattt atataattat aatacattat ataatttata 660
cattataatt atat
                                                                 674
<210> 459
<211> 682
<212> DNA
<213> Homo sapiens
```

```
<400> 459
tttttttaaa tccatggctt gttaattgtc atcccagtta tttacatgtg actatagaga 60
ctgcattctc ccagctgcca ggccgccagg gctttgccac tggtataatt tataacacga 120
ctaattaaaa tgaatttgct tgcaataagg ttctgtgtgc tatttgtggg agaggagtta 180
ttaaaatttt caqtacagta atagtaaact tgaatgcaaa gtaataataa tcatacattt 240
ttaattacat gtttaatacc catttggcta atgtagaact attctgaaaa ttacttggga 300
tcagcacaat gtctttttgt gcttagtagt atccaaagac atccttctga atgggcttag 360
caatatqcac tgtcatcaag atacagctgt ttgatgacag acacacagtg tgttcctatg 420
atactttgca caagatcagc tatgacaaat acaagttcat tttgcttatt gcaggcaaat 480
aatgtccttt gcaggaactt ggatggagcc agaggccatt attctaagtg aaatacctca 540
qqaqtqqaaa accaaatacc atatgttctc acttacaagt gggaactaag ctatgggtac 600
acaaacgcat atagagtaat ggactctggc gactcatact acatattgag tacaatgtac 660
actacttggg tgatgggtgc ac
<210> 460
<211> 663
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(663)
<223> n = A, T, C or G
<400> 460
gaattegaac ceettegegg ggegegegag eggegeeage teggggeage ggaaceeaga 60
gaagetgagg gggeggtage ggeggegaeg gegaegaega egaeteeege gegtgtgeee 120
agectettee egeogeagee gecettttee teecteeett aegteeeega gtgeggeagt 180
accgcctcct teccageege geggetteet ecagacetet eggegegggt gageeetatt 240
cccagaggca ggtggtgctg accctgtaac ccaaaggagg aaacagctgg ctaagctcat 300
cattgttact ggtgggcacc atgtccttga agcttcaggc aagcaatgta accaacaaga 360
atqaccccaa qtccatcaac tctcgagtct tcattggaaa cctcaacaca gctctggtga 420
agaaatcaga tgtggagacc atcttctcta agtatggccg tgtggccggc tgttctgtgc 480
acaagggcta tgcctttgtt cagtactcca atgagcgcca tgcccgggca gctgtgctgg 540
gagagaatgg gcgggtgctg gccgggcaga ccctggacat caacatggct ggagagccta 600
agcctgacag acccaagggg ctaaaganaa gcagcatctg gcatatacag gctcttcgac 660
                                                                   663
tac
<210> 461
<211> 612
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(612)
<223> n = A, T, C or G
<400> 461
ttttttggga tccaatctnt ttattgtcag ggtcccctcc ctgnggcccc ccgccaaacc 60
tatagaaaaa acccaagcct gggagtgtcc tggggagggg aggtagtatg gggaaacccc 120
tgtgctctac cctntggcct gggcagtgca nacagggagg gctcatgggg aaggagtagg 180
ccagtaactc cacctgcana ggacatggca ctggctggga tgcgttgggg gaggaggcgc 240
ctgctgccag ctttcctntg gtacccgctg gggggtggca tccagggttg ggtgcccggc 300
```

```
ttgaggcctg gggcagcgat gcccttcacc tgctggnggc cattgctcct gtcaggctgc 360
ttactgcaag gccccatcat ccgcgtctgt gtcctggctg tgttccagct cttcctcgct 420
gngtgtcagg agcccttcct catcgccgtc gtctcgggtc cgtgcttccc cctggggcag 480
caccaccact ggcaccggca ccgntgcacc accaccgccg ccgccgccgn tggngccacc 600
ttcatcaccc tt
<210> 462
<211> 672
<212> DNA
<213> Homo sapiens
<400> 462
gaattcgaac cccttcggat ggaaggggcc ggggcagcgt cggggaaagg aagggccgga 60
ggcgcggcgg cggcggccg agaggggcgg cggcggcggc ggcggcgggg ttcccgcgcc 120
geggageeeg geeegagage egegteeaeg tteetgeete etgeteeege egeeetgggg 180
egeogecatg aegeoegate tgeteaactt eageoecaga tgteaceaag eteteggaet 240
ctaacaagga gaacgcgctg cacagctaca gcacccagaa gggccccctg aaggcagggg 300
agcageggge gggetetgag gteateagee ggggtggeee teggaaggeg gaegggeage 360
gtcaggcctt ggactacgtg gagctctcgc cgctgaccca ggcttccccg cagcgggccc 420
gcaccccage cegeacteet gacegeeetg gccaageagg aggagetgga gegggaeetg 480
geccageget eegaggageg gegeaagtgg tttgaggeea cagacageag gaceccagag 540
gtgcctgctg gtgaggggcc gcgccggggc ctgggtgccc cctgactgag gaccagcaaa 600
accggcttag tgaggagatc gagaagaagt ggcaggagct ggagaagctt gcccttgcgg 660
gagaataacc gg
<210> 463
<211> 562
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(562)
<223> n = A, T, C or G
<400> 463
ttttttaaag tataaagtgt tttggaaaaa aaggaaaaan ntctatataa aaatctcttc 60
acatataaaa tootgaagaa ggtgcaaggt gagacccagt gcgaggggcg tgctcagata 120
tgcagtgtgt gtgtgtgtgt gtgtgtgtgt gtatccgtgt gtacatgtgt gcacgtgtgt 180
agtgcacgtg tggcccacag agggtgggga gaaagcttgg ctttttactt ccatccagga 300
gggaaggagg gcggctggtc ctccagcctg gagggtctgc agctgggcgg gacctctact 360
cagccagget gttgcgcatc gactcettet cetggaggge ggccatggca agacgcaggt 420
geteetteag etgetegate teeegeteag acceptett gatgtggete aacteeacat 480
agacgteetg gtacttteee naggtgaage gettgteett etgeateate tggagetegt 540
                                                              562
cccggaggca ctgcaccttc ct
<210> 464
<211> 553
<212> DNA
<213> Homo sapiens
<400> 464
```

```
qaattcqaac cccttcqqqa ccaqqaaccc aggaqaqcat ggccacqctg cgccqgcttc 60
gggaggcgcc gcggcactta ctggtttgcg agaaatccaa cttcggcaac cacaagtcgc 120
gccaccggca tcttgtgcag acgcactact ataactacag ggtttcattt ctcattcctg 180
aatgtgggat actatcggaa gaactgaaaa acctggtcat gaacactgga ccctattact 240
ttgtgaagaa tttacctctt catgaattaa ttacacctga attcatcagt acctttataa 300
agaaaggttc ttgctatgca ctaacataca atacacatat tgatgaagat aatactgttg 360
ccctgctacc aaatgggaaa ttaattttgt cactggataa agacacttat gaagaaactg 420
gacttcaggg tcatccatct cagttttctg gcagaaaaat tatgaaattt agttcagaag 480
aatcgacaat gatgtcatat ttttccaagt accaaattca ggagcatcag ccaaaagtag 540
                                                                                                                                 553
cactgagccc gtt
<210> 465
<211> 383
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(383)
<223> n = A, T, C or G
<400> 465
tttttggaag aaaacacgat ttttaatttt tatttttat gggggacagn gatcatttgc 60
cccaacagcc atntgaagcc aatagtcctg attattaaaa atcacaaagt tatataaatg 120
ntctcctcct tttcgaaaac catgttcatt tttttcccaa naaacagggc tgtctgcaaa 180
gccttgaacg gacagngtaa cccatggagc taacttcggt tcatcaaagt agngacagan 240
atgttccaat agganacaga tcttntntgg aagtatgaag ccagngattg tacacaaata 300
agcttttgcc accactgtgc ttggctcagg acagcaatag gttgatatga aattattagg 360
ctcattattt aggncgacat tac
                                                                                                                                 383
<210> 466
<211> 673
<212> DNA
<213> Homo sapiens
<400> 466
gaattegaac ceettegete eeteetgeac geaatggtgg eetatgatee egatgagaga 60
ategeegee accaggeet geageacee tacttecaag aacagaggaa aacagagaag 120
egggetetgg geagecacag aaaagetgge ttteeggage accetgtgge acceggaacca 180
ctcagtaaca gctgccagat ttccaaggag ggcagaaagc agaaacagtc cctaaagcaa 240
gaggaggacc gtcccaagag acgaggaccg gcctatgtca tggaactgcc caaactaaag 300
ctttcgggag tggtcagact gtcgtcttac tccagcccca cgctgcagtc cgtgcttgga 360
tctggaacaa atggaagagt gccggtgctg agacccttga agtgcatccc tgcgagcaag 420
aaggtagege ggaaccaget tetetgaegg egetgetett egacecagee eaggeegeea 480
ctgaattttg tgtctgtaat ttttctttga cagacagatc cgcagaagga ccttaagcct 540
gccccgcagc agtgtcgcct gcccaccata gtgcggaaag gcggaagata actgagcagc 600
acceptcettcega general agencegacce general gen
ctgctgagac gcc
<210> 467
<211> 373
<212> DNA
<213> Homo sapiens
```

```
<220>
<221> misc_feature
<222> (1)...(373)
<223> n = A, T, C or G
<400> 467
tttttactgg aacgacagct tattntttaa taaaagtcag gggngtcagc agngtcactg 60
gtaanacatg atggcgctcc acgactgacc agcagcgctg ggaagggaca cgcanaaccc 120
accttccaac cacgcccaac acatnacana aatgcctgct cgtttgtttt gattcatata 180
caaagttaca aagtatttcc tgccccaaat tnttaacgaa aatgaaagaa aaccctanaa 240
tgcgggggtt ttacaagtat attagcccan aacatcctag gcagctgcnc gggccgcggg 300
tgcggcaggg cgcagggcaa cacccaaagc cccggccagc gcgaaacgga cgcaggcgca 360
tccccagccc tcc
<210> 468
<211> 573
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(573)
<223> n = A, T, C or G
<400> 468
gaattegaae eeettegetg etgteetaet tgatgettgt eaetgteatg atgtggeeee 60
tngctgtgta ccaccgactg tgggatcgag catatgtgcg gctgaagcca gctctgcagc 120
ggctagactt cagtgtccgt ggctacatga tgtccaagca gagagagaga caattacgcc 180
gcagagetet ecacecagaa egageeatgg acaaceaeag tgacagegaa gaggagettg 240
ctgccttctg tcctcagctg gacgattcta ctgttgccag ggaattggcc atcacagact 300
ctgagcactc agacgctgaa gtctcctgta cagacaatgg cacattcaat ctttcaaggg 360
gccaaacacc tctaacggaa ggctctgaag acctagatgg tcacagtgat ccagaggaat 420
cetttgccag agacetteca gacttecett ceattaatat ggateetget ggeetggatg 480
atgangacga cactagcatt ggcatgccca gcttgatgta ccgttctccg ccagggggct 540
                                                                   573
gaggagcccc aaggccccac ctgccagccc ggg
<210> 469
<211> 635
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(635)
<223> n = A, T, C or G
<400> 469
tenegateta gaactaggtt ggacaggett geteaagttt caccagagtt antactggee 60
tctgttcgca gagtttttag ttnnacactg cagaattggc agactacacg gtttatggaa 120
gttgaagtag caataagatt gctgtatatg ttggcagaag ctcttccagt atctcatggt 180
gctcacttct caggtgatgt ttcaaaagct agtgctttgc aggatatgat gcgaactgta 240
agtatactgg agataatttt gaccataaat ttctgttttc agtataagct aatgggagtt 300
ccttaattgt tagagcttag tatatgttaa taccggggca ttttgatgtt gcaataaata 360
agaagaggtt teetaaettt tteetgatet agetggtaae ateaggagte agtteetate 420
```

```
agcatacatc tgtgacattg gagttcttcg aaactgttgt tagatatgaa aagtttttca 480
cagttgaacc tcagcacatt ccatgtgtac taatggcttt cttagatcac agaggtctgc 540
ggcattccag ngcaaaagtt cggagcagga cggcttacct gttttctaga tttgtcaaat 600
ctctcaataa gcaaatgaat cctttccttg aggat
                                                                   635
<210> 470
<211> 593
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(593)
<223> n = A, T, C or G
<400> 470
qaattcqaac ccttcqqtat taacaaatat ntacatttct atttttataa tccataaqqa 60
tatgcctgtt ttaaataaca tacatattaa caatatctat caggaaaacc ctcaagacag 120
cttctagtta aaaccttngn tgctgtcctc tcaaactata tttataaaaa tttgctaggg 180
ccaaatccat acttgcagaa taattcatca aattttattt ttaagngaaa agtaaccttt 240
caggicattic agcagicatac attigacaatic taggigtatat atgitatigtat gitticitatt 300
gtatgtctat atatgtatgt ggggaggaca ggagtgaatg ttcacacact tttcttgcgt 360
actcaactaa attggagaat gtttctgaag aaaattggat gaaattagct gctgagattg 420
agtttctgcc ttaaaatctg aaacaaaaaa agggacaaat tgctggtang atctactgac 480
tqtnqccatc accaqaacac ttagtttctt cccagacatg aatttcctga caggctctga 540
gccagaaaca cactgtgggc gtgcatntgg gtcaccctgg atatgcctcc act
<210> 471
<211> 581
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(581)
<223> n = A, T, C or G
<400> 471
tttttttaat cangggacat ttattaacat gcttcaaaag tgaccaaagt gtccagccag 60
cacaatagcc gaggcaatca acqttctctt agtgtgtgat ctcgtccaaa acaccaaata 120
aataqqttta qqaataacct caaataaatt qtaatttaac ttcqcccaaa attatacatc 180
ctctactgct cttccctgct cctgtaaaga tactagcggg aggggagaaa gctcaaatga 240
ctctqtaatt taqaattaca accaqaqaaq aaatacttca aqcacaataa aqacqttcca 300
ttgaagagcg acattcattc tggaatgttt gttttgaaaa caactcttnt gggggaattc 360
aaaaggtact gaacaaagca acataaagta agttttgggt tgttttgcaa aataaaaata 420
tacaattgag tggaccagat ggcaaaaaca taccaattac aatctgaatg ctatatttaa 480
aacccttaaa ttctgaaggc ctgaatatca acaaacctat ttatgtttat gatcctaaaa 540
agacattaaa tattattaaa cccccaactt ccaaaacata q
<210> 472
<211> 674
<212> DNA
<213> Homo sapiens
```

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<220>
<221> misc_feature
<222> (1)...(674)
<223> n = A, T, C or G
<400> 472
gaattcgaac cccttcggat ggcgtgatgt ntcacagaaa gttctccgct cccagacatg 60
ggtccctcgg cttcctgcct cggaagcgca gcagcaggca tcgtgggaag gtgaagagct 120
tecetaagga tgaecegtee aageeggtee aceteacage etteetggga tacaaggetg 180
gcatgactca catcgtgcgg gaagtcgaca ggccgggatc caaggtgaac aagaaggagg 240
tggtggaggc tgtgaccatt gtagagacac cacccatggt ggttgtgggc attgtgggct 300
acgtggaaac ccctcgaggc ctccggacct tcaagactgt ctttgctgag cacatcagtg 360
atgaatgcaa gaggcgtttc tataagaatt ggcataaatc taagaagaag gcctttacca 420
agtactgcaa gaaatggcag gatgaggatg gcaagaagca gctggagaag gacttcagca 480
gcatgaagaa gtactgccaa gtcatccgtg tcattgccca cacccagatg cgcctgcttc 540
ctctgcgcca gaagaagccc acctgatgga gatccaggtg aacggaggca ctgtggccga 600
gaagctggac tgggccccgc gagangcttg agcacaggta cctgtgaacc aagtgtttgg 660
gcaggatgaa aatg
<210> 473
<211> 646
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(646)
<223> n = A, T, C or G
<400> 473
ttttttcagn ggaaaataac ttttattgan accccaccaa ctgcaaaatc tgttcctggc 60
attaagetee tinticetit geaatteggi ettietteag nggiceeatg aatgetitet 120
tctcctccat ggtctggaag cggccatggc caaacttgga ggnggtgtca atgaacttaa 180
ggtcaatctt ctccanagcc cgccgnttcg tctgcaccag caaggacttg cggagggtga 240
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caccatagng gacaaagcca cccanagggt tgatgctctt gtcanatagg tcatagtcag 360
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aaatcttctt gttgatctca gtgcggtgat ggtagccttt ctgcccagcg cgtgccacag 480
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gggtcttgcg gggcagcttc ttggtgtgcc aacgactggt gacccctttg tagcctttgc 600
ccttggtcac cccgatgacg tcgatcatct catcctgccc aaacac
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<210> 474
<211> 544
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(544)
<223> n = A,T,C or G
<400> 474
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gcagcacggc acagcacgct cgacttcatg ctcggcgcca aagctgatgg tgagaccatt 180
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caggaccatg gctatttagc aacctacaca aacaagaacg gcagctttgc caatttgaga 300
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aaagaagaga tcgacagtat tttgaacaaa gtagaggaaa gaatgaaaga attgagtcag 420
gacaagtact gggcgggtga aacgattacc acccatagtg cgaggaggag ccatcgacag 480
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gacg
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<210> 475
<211> 578
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(578)
<223> n = A, T, C or G
<400> 475
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ctcacagggc agacccctgt gttttccaaa gctagataca ctgtcagatc ctttggcatc 180
cggagaaatg aaaagattgc tgtccactgc acagttcgag gggccaaggc agaagaaatc 240
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gacaagaagc gcaggacagg ctgcattggg gccaaacaca gaatcagcaa agaggaggcc 480
atgcgctggt tccagcagaa gtatgatggg atcatccttc ctggcaaata aattcccgtt 540
tctatccaaa agagcaataa aaagttttca gtgaaaaa
                                                                   578
<210> 476
<211> 619
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(619)
<223> n = A, T, C \text{ or } G
<400> 476
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tettececeg taaggaaatg geeggggage teeaggggae eeaggegeeg tegettegge 120
ggagcctggg ctgaccagcc aggacagcgg ggtaaacccg aacaattctg cgcgaggtag 180
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tgcaatgaaa tctcgaaggg gacctcatgt ccctgtggga cacaatgccc ccaaggactt 360
gaaagaggag attgatattc gactctccag ggttcaggat atcaagtatg agccccagct 420
ccttgcagat gatgatgcta gactactaca actggaaacc cagggaaatc aaagttgcta 480
caactatctg tataggatga aagctctgga tgccattcgt acctctgaga tcccatttca 540
ttctgaaggc cggcatcccc gttccttaat gggcaagaat tttccgcttc taccttgctg 600
gatcttgcga aacactagt
                                                                   619
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<210> 477
<211> 674
<212> DNA
<213> Homo sapiens
<400> 477
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aaggaacttg tttcttcaag ctcttctggc agtgattctg acagtgaggt tgacaaaaag 120
ttaaagagga aaaagcaagt tgctccagaa aaacctgtaa agaaacaaaa gacaggtgag 180
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cagattggga aaatgaggta cgttagtgtt cgcgatttta aaggcaaagt gctaattgat 300
attagagaat attggatgga teetgaaggt gaaatgaaae eaggaagaaa aggtatttet 360
ttaaatccag aacaatggag ccagctgaag gaacagattt ctgacattga tgatgcagta 420
agaaaactgt aaaattcgag ccatataaat aaaacctgta ctgttctagt tgttttaatc 480
tgtcttttta cattggcttt tgttttctaa atgttctcca agctattgta tgtttggatt 540
gcagaagaat ttgtaagatg aatacttttt tttaatgtgc attattaaaa atattgagtg 600
aagctaattg tcaactttat taaggattac tttgtctgcc cacccctagt gtaaaataaa 660
atcaagtaat acat
                                                                 674
<210> 478
<211> 663
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(663)
<223> n = A, T, C or G
<400> 478
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acagacatca teccatggtg aacatgttta ataagtgaaa geaagteaga cateteatet 120
aagtcattat tttctgcaga ctaagcaata actacacaga acactatggg taaacaaaca 180
cctgctcagt tttcacacaa gccatgttgt ttatcaaatt agatctgcta atattgaata 240
cagtagattc ggtgattgta gttctcatat aagtatctta ttgagataac attttgacag 300
tttcactgac tttccaaata agcataccat aatcaaagaa aagaataaag agtgaagtaa 360
aaccattggg ggtggaagtc aaacaagcct agacatttga ttggaagaga aaagatcaaa 480
tatgaagttc acaaaccaaa agtttataaa ctcaatgcaa tacaaatcct ttttattgta 540
aaagctgagt tgaaactaaa agatctataa aaactgttac ttttggcctt aaacagtacc 600
aactettatg atcaaaaaag gecacaeagt taagattgna ttaettgatt ttattttaca 660
cta
                                                                 663
<210> 479
<211> 673
<212> DNA
<213> Homo sapiens
<400> 479
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teettaacaa ggaaateace agtgtgggea gtteeaageg ggaagaggag geeaagteag 180
aacttcgagc agccattgat cggtatgtgc aagagaagat tgtgctagca gctcaggcaa 240
```

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tttcacgctt tgcttaccag aagatcagta atggagatgt gatcctggta tatggatgct 300
catctctggt atcacgaatt cttcaggagg cttggacaga gggccggcgg tttcgggtgg 360
tagtggtgga cagccggcca tggctggaag gaaggcacac actacgttct ctagtccatg 420
ctggtgtccc agcctcctac ctgctgattc ctgcagcctc ctatgtgctc ccagaggttt 480
ccaaggtgct attgggagct catgcactct tggccaacgg gtctgtgatg tcacgggtag 540
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aaacatacaa gttctgtgag cgtgtgcaga ctgatgcctt ttgtctctaa tgagctagat 660
gaccctgatg atc
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<210> 480
<211> 203
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(203)
<223> n = A, T, C or G
<400> 480
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gcccagggag tagtggaggc cgntggcagc ttcggggctt atggtgccca ggaggaagcc 180
cantgcccta ctctgcattt cct
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<210> 481
<211> 482
<212> DNA
<213> Homo sapiens
<400> 481
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caaccagett gtagattttc agtggaaact gggtatggct gtgagetcag acacttgcag 120
atctcttaag tatccttacg ttgcagtgat gctaaaagtg gcagatcatt caggccaagt 180
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ggaaattgct gcagttattg aaacggtgtg aagacggatt ctttggttga taaattgcta 300
tcattctaaa gtcatggact tcactttcgg caacaaaact aaataaggat ggaacattta 360
ttgaatgaaa aatgcacttt tgtttttcca tttttttaaa taataaaaat cagacaaaca 420
gaaaaaaaaa aaaaaaaggg cggccgctcg agtctagagg gcccgtttaa acccgctgat 480
<210> 482
<211> 505
<212> DNA
<213> Homo sapiens
<400> 482
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cctaggtgat aaggtgactg gacccagtaa accctttgtg tgctgggggg ttttatgcct 120
tgtagaaccc agtgtgagca agatttgggt accctacata cattcagtag ccaggaaagg 180
gtgattggat tgccagactc tgcctgctgg caaaaggatg agctgtagaa gctgaagtcc 240
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atttggaatt gaatttttcc tctaattatt ctagggaaac cctgggctaa gaaaccaatg 360
taaaacctga tgaggtagtc tgtagtcaca ctgggtagag gtagaggcaa ccacaaaatt 420
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tggtaaaaat ttaggtcgga tgcag
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<210> 483
<211> 501
<212> DNA
<213> Homo sapiens
<400> 483
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ttcaagtggg caagctgggg tttttaggta gtcagtggcc tagttcctaa agccacagta 180
taggatetgt taaactgaat gtetgttgaa agtttgtttt agetgettgg aggetteett 240
ttaagacaaa ctgtatgtga ttaagttgtt ttgagggaac tgaagaacct gatgtagccc 300
ctggccagat aactgcctga tttctcagat attatttctc tgggaaacat tctacatagc 360
acaggagett aagagtggea ttatettete geettaattt eeagagatta tttetgtaet 420
qaqaatcctg gaactactat gctaggaaat ttaaagctgc atggtctgtc ttgttttcat 480
ttaattattg tgaataccta g
<210> 484
<211> 501
<212> DNA
<213> Homo sapiens
<400> 484
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aaagggtgct acccaagccc ctcttacctc tctggatgct ttctttaaca ctaactcacc 480
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                                                                   501
<210> 485
<211> 504
<212> DNA
<213> Homo sapiens
<400> 485
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cattatatat catataaaat aaacctttaa atattgaaat gaaaagataa aaatacatac 240
actaagtgaa taggtcaaaa gtgtgagatc atcttgaaca ttatcttgaa gagaagatac 300
caatttacct tetgeteaga teatggtgta egatateaca acetgeetag aataaetete 360
cttttctgaa ccatttattc actacttttg tcttccaatt aaatattagc ctgacttcaa 420
atatcataca ttagtttcct ttgtttatgt aattgaatta tataacatat attcattaga 480
gcctattttt tttaaaattt ttgt
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<210> 486
<211> 501
<212> DNA
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<213> Homo sapiens
<400> 486
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gcacggacgc accaaggagc agaaggggcc cctgtcgggt gcagcgtcct gggagcatat 240
caaggetgtg eggaaggetg tggecateee tgtgtttget aaegggaaca teeagtgeet 300
gcaggacgtg gagcgctgcc tccgggacac gggtgtgcag ggcgtcatga gcgcagaggg 360
caacctgcac aaccccgccc tgttcgaggg ccggagccct gccqtqtqqq aqctqqccqa 420
ggagtatetg gacategtge gggageacce etgeceeetg tectaegtee gggeeeacet 480
cttcaagctg tggcaccaca c
<210> 487
<211> 501
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(501)
<223> n = A, T, C or G
<400> 487
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caacccctat aaagggtagt tottttaaaa aaaatttott tattggcaac aacataaaag 180
atatgaaaga atcactcata atttatcagc ataacatagc tattctcatt tttgcaattg 240
actttttagt tcttgaccaa atgtaatttt tattagttgt gattaactga ttttgtgctt 300
tttttaaaaa aaaaaaaaac ctagaataag acatttgttt tgttaattat tataaatgac 360
tgtattcatt ctgtttatgt accataattt tggatgttcc tacgatgtta aacttttagg 420
ttgtttttaa ttgtttgttc ttatagacaa ctctgtaagg gnttttaact gcttttatca 480
ggagaatgtc aaagaagtcc t
                                                                   501
<210> 488
<211> 148
<212> DNA
<213> Homo sapiens
<400> 488
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ggacagaggt ggaattgagt gtccacaggc cagctgagga ggtggtaccc agcactctat 120
gaaccettcg ctcaagtcag cctggagt
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<210> 489
<211> 501
<212> DNA
<213> Homo sapiens
<400> 489
gctgtggatt cccctccaag tggaggagga tgggcaggct ggggatcctg gggcaaatct 60
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ggagccactc tacggattca tggtgtaaat tctggatctt ctgaaggagc ccaaccaaat 180
actgaaaacg gagtccctga aataacagat gcagccacag atcagggccc tgcagaaagc 240
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ccacccactt ccccttcatc agcctctcgg ggtatgctgt ctgccatcac caatgtggtt 300
caaaacacag gtaaaagtgt cttaactgga ggccttgatg cgttggaatt catcggcaag 360
aaaaccatga atgtccttgc agaaagtgac ccgggcttta agcggaccaa gacgctcatg 420
gagagaactg tttccttgtc tcagatgtta agggaagcta aggagaagga gaagcagaga 480
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                                                                   501
<210> 490
<211> 482
<212> DNA
<213> Homo sapiens
<400> 490
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gcagtcactc aggtttattt ccaccagggc ccaagaaaaa aagaaatgag gcaacctaaa 360
attocatcaa gatagatacc aatatccaag gtgcttggtc ttagcggtgt gggacccacg 420
ttaaggetet tggtgggaag gtgggaggtg ttttcageat gagatagggt tcaggetgtg 480
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<210> 491
<211> 483
<212> DNA
<213> Homo sapiens
<400> 491
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ggggaaatgg ggctgggggc cgtccccggg agacaggcgg ccttccgaga gggactggag 480
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cag
<210> 492
<211> 266
<212> DNA
<213> Homo sapiens
<400> 492
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gaaggeeeta gatgtgggge ttetagatta eeceeteete etgeeataee egeacatgae 120
aatggaccaa atgtgccaca cgctcgctct tttttacacc cagtgcctct gactctgtcc 180
ccatgggctg gtctccaaag ctctttccat tgcccaggga gggaaggttc tgagcaataa 240
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agtttcttag atcaatcaaa aaaaaa
<210> 493
<211> 483
<212> DNA
<213> Homo sapiens
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<400> 493
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cgcggaagaa gaagggcccg gggcccctgg ccacggcgta cctggtcatc tacaatgtgg 180
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ccttattgga gattttacat tgtgctatag gaattgttcc atcttctgtt gtcctgactt 360
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tacagagtga agacagtgtc ctcctgtttg ttattgcatg gacgatcacg gaaatcatcc 480
gtt
                                                                 483
<210> 494
<211> 301
<212> DNA
<213> Homo sapiens
<400> 494
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ttctgaagct gggtggttca cacctgtaat cccagcactt tgggaggatc tcttgagccc 180
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<210> 495
<211> 496
<212> DNA
<213> Homo sapiens
<400> 495
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cgtgggaggg cgcgagcgaa cgcgggcgag gagcggccga gccgctgaag aggagctggg 180
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gggattette geceaaaatg gagttaatee tgaetgggag aagaaagtaa ttgagtattt 360
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tgaccctgag ttttac
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<210> 496
<211> 494
<212> DNA
<213> Homo sapiens
<400> 496
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ctttctatac ctttgtacta tgcactgccc tattgattct acacccaata atgatattac 180
ttgaacccat ctgtaagaaa ctgcttcgga aattcatttg tgtgtatgta aataacacaa 240
catagaaaca ggaagggaaa aaagtctgca gtaatgcacg tattttttt ctttcctgtt 300
tattttcggt tttgctttaa gtccttttat ttttaattcc ctttttgttt ttctttttgg 360
gttttggttc cttttgggtt tatgggtgcc ctgatactcc agcagagatc agaaggctac 420
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<210> 497
<211> 184
<212> DNA
<213> Homo sapiens
<400> 497
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gtcg
<210> 498
<211> 471
<212> DNA
<213> Homo sapiens
<400> 498
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atacttagtg atatactttg cttgaaaatc actcagcaaa gtagttcaca tgatgtgtat 120
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<211> 101
<212> DNA
<213> Homo sapiens
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<212> DNA
<213> Homo sapiens
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tcgggaaaaa ctagccaaaa tgtacaagac cacaccggat gtcatctttg tatttggatt 180
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<210> 514
<211> 346
<212> DNA
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accaactttt caaacatcca ggacaaccag ttttctccct gtggtgtgcc catttcgctt 180
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ggacctatgc cggggacagc actggaagca gggtacagta gcccaaagaa aaagacacat 300
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<210> 515
<211> 549
<212> DNA
<213> Homo sapiens
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teccagaate ageagacate aaateeaaeg cacagtteag aagatgtgaa geeaaaaaee 120
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<211> 565
<212> DNA
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<211> 127
<212> DNA
<213> Homo sapiens
<400> 521
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127
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<210> 522
<211> 642
<212> DNA
<213> Homo sapiens
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<223> n = A, T, C or G
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<211> 631
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(631)
<223> n = A, T, C or G
<400> 529
acttgcctaa agtttttata tctgnntctt ctgctgtaaa tcttcccttc ataaatgaaa 60
attttaataa aatcaactat gtggaaatat ataattaaag gaattcacta actgtgattt 120
tcataattta gggacattct cttctagtaa gcatggtgca ttatttacta gagatataat 180
atgcattaaa acaaaaaatg ttttctatca tcatagaaaa gtttgaggtc cagggataat 240
catctctgga tacattattt cctaccgtcg tggtacacac tgaacacatt tgaggcttat 300
gactggttct tttacttaca aatattgttt agacacattt tcaaatgtca caccaatcaa 360
taataataag gaatggattt tatctatatt gacagttctt tcaaccttaa gagtgaactg 420
ctacaggtaa gattcaatca catttttcag gagaaagcta ttgagaccaa tatgctttgg 480
ttatctaata ggggtggaat gacttataat gctatttact ccaggcaaag agaaaataca 540
acagacatag gatcttgatt tcaacgtagt tctcctccat gtgcatttct ctgtccgttt 600
aggcaatgcc aactggtcca ccagtgaaca t
<210> 530
<211> 316
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(316)
<223> n = A, T, C or G
<400> 530
acacatttaa atgactcacg agantnaagt ttttttcaaa tatattaaga tcacaccacc 60
ttgttgttta tcgaaagata ttcaaggaga aagatctgac tctccaaact gcatctgaga 120
ttgccacttt aaacagacct catttcaaac atgcaacaac gccactggta ataaagcttt 180
ggaatgggtg ctcattctat tatttcacta caaacagcat agaaagcaag agaagttggg 240
aatttattet aaaatagaat ggaggttgte atetacagea geacteetea eteetetgtt 300
                                                                   316
gccattttta gcaagt
<210> 531
<211> 296
<212> DNA
<213> Homo sapiens
<220>
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<221> misc feature
<222> (1)...(296)
<223> n = A, T, C or G
<400> 531
aaagtatcat ttatttgaaa aacatacatt atcattntgt ttttgatatt tgataatgaa 60
aaaaatcttt gnttgtttat ttctgaaaaa gaactgtatt tagngattat tttagatagt 120
gatattatan cattcatctg tgtgtaaatt atttcatata gggaagagtt ctgatctgta 180
cctatggttc ttattgaaaa caacattgga tgtgcatttc tgtgatgtta tgaatacatt 240
tctactttat tttgaaacat ttgccaaact aaatactgta acactgtata acattt
<210> 532
<211> 266
<212> DNA
<213> Homo sapiens
<400> 532
acatatgcac caaattccat tttagaagtt tccatatcat tttcatagaa aacaaagttt 60
gaaaacaagt aacatttaaa cacagcacgg tattctacca caactgaaac ttttttcttc 120
ttcttcttta caggactcaa caaaatctaa aaatgaacta tgctgtagat ttacctcatg 180
caaagatctt tatgttatct ctgaaaatga aaaggatggc cttttaagca cattttactg 240
                                                                   266
ttttatacta ttatggcaac ttgtgt
<210> 533
<211> 289
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(289)
<223> n = A, T, C or G
<400> 533
actcagaagt cacttttaat atcancgaca gaaatatttc actaattcaa ctgaggcaaa 60
tttcctttct agacaaagga cctagaaatt gagcatgcaa aacatccatc cattcattca 120
ttcaaataat tagccaattt taccgtcatt taattccacc agaagcaaat actagaatat 180
ctagaagtag tttgggtaaa gaaacattta cattttaata ttgtgtaatg tcataaattt 240
ggggctaaaa taacaccagg tcaaatttga tccctttgta tgtgagggt
<210> 534
<211> 293
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(293)
<223> n = A, T, C or G
<400> 534
aaaataaaag gttctttaca agatgatacc ttaattacac tcccgcaaca cagccattat 60
tttattgtct anctccagtt atctgtattt tatgtaatgt aattgacagg atggctgctg 120
cagaatgctg gttgacacag ggattattat actgctattt ttccctgaat ttttttcctt 180
```

```
tgaattccaa ctgtggacct tttatatgtg ccttcacttt agctgtttgc cttaatctct 240
acagcettge teteeggggn ggttaataaa atgcaacaet tggcattttt atg
<210> 535
<211> 408
<212> DNA
<213> Homo sapiens
<400> 535
acttgaacac ttaaagagaa aaactctaaa taaagtcata gaggggatgg tagagatgac 60
cacagaaaat gaccacggag agtattatga agattgcaag attagacatt gatgatgtaa 120
attactccct ttctagataa aataatccat agatgtttat gaatcatatt tgtatgatta 180
ttgctgttac tattattttg acacattatt tattattatt gttgtcacta ttattaccat 240
taagatagca ggcgtaaaac tgtactggtt ccttcagtag tgagtatttc tcatagtgca 300
gctttattta tctccaggat gtttttgtgg ctgtatttga ttgatatgtg cttcttctga 360
ttcttgctaa tttccaacca tattgaataa atgtgatcaa gacaaaaa
<210> 536
<211> 184
<212> DNA
<213> Homo sapiens
<400> 536
acctctcatc aaggetetge ctacaggeac attgtgatgt atetetgeac tgatcaccta 60
ggtcatgtaa cttttttcta ggctctacct acgatggcat tgtgacataa ctctgcacta 120
atcatccacg tgatgtaact cttgtctagg atgtgcctaa attaactttt tgacgtaacc 180
ctgt
<210> 537
<211> 311
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(311)
<223> n = A, T, C or G
 <400> 537
ccacagttgt atcatatagc atctntaaca tttcatctag gattatctag tatagatctt 60
actatatttg gggctatgtt gtatacaatg ttaacaagaa catatettet etgcatatat 120
gtgtgaatta taaagaaaag catgagaatg actctaagtt caacaaacat gggtgaatct 180
 ctatgtgctc ccagtgtcct ggatgggctc cccagcaagc cattcctcct tcctgttctg 240
 atattactat tcttttttac attgtgctaa ggaggacaaa aggtgagaga tgaaaataaa 300
                                                                    311
 gccttgcctt t
 <210> 538
 <211> 302
 <212> DNA
 <213> Homo sapiens
 <400> 538
 aaaataaaaa agcaaaaact cttgtggtac ctagtcagat ggtagacgag ctgtctgctg 60
 ccgcaggagc acctctatac aggacttaga agtagtatgt tattcctggt taagcaggca 120
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<210> 542

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ttgctttgcc ctggagcagc tattttaagc catctcagat tctgtctaaa ggggtttttt 180
gggaagacgt tttctttatc gccctgagaa gatctacccc agggagaatc tgagacatct 240
tgcctacttt tctttattag ctttctcctc attcatttct tttatacctt tcctttttgg 300
                                                                   302
<210> 539
<211> 396
<212> DNA
<213> Homo sapiens
<400> 539
actgtttatt tgctccttct cttcatgcct gtggctggat gtcccacaac actataagaa 60
atataagtca agccctttgt gttaagcaag aactacagac tccatctttt cacccaaatc 120
atgaatgacc aataaaaagc aagttattcc agaggaagaa gcagcccttg aaatgttaag 180
gcttaggctt gaaaggtgaa gagcaggaat tctctctttc aaatcctaga gcataaaccc 240
atgtgtggcc aagtgagatc agccctcaag ggcacatgcc aagggcagag cagcccatgt 300
agacagette ggagggeatg ggggtgtagg gagttegggg tageteetea ttaactattt 360
gttgggtgag taaaggggtg aggctcagtg gcaggt
<210> 540
<211> 634
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(634)
<223> n = A,T,C or G
<400> 540
ccaaaaacaa gatgaccaga tttgntttna gcctgatgac cctacaggtc gtgctatgat 60
atggagteet catgggtaaa geaggaagag agtgggaaag agaaceaeee caetetgtet 120
tcatatttgc atttcatgtt taacctccgg ctggaaatag aaagcattcc cttagagatg 180
aggataaaag aaagtttcag attcaacagg gggaagaaaa tggagattta atcctaaaac 240
tgtgacttgg ggaggtcagt catttacagt tagtcctgtg tctttcgact tctgtgatta 300
ttaaccccac tcactaccct gtttcagatg catttggaat accaaagatt aaatccttga 360
cataagatct catttgcaga aagcagatta aagaccatca gaaggaaatt atttaggttg 420
taatgcacag gcaactgtga gaaactgttg tgccaaaaat agaattcctt ctagtttttc 480
ttgttctcat ttgaaaggag aaaattccac tttgtttagc atttcaagct tttatgtatc 540
cateceatet aaaaactett caaacteeae ttgtteagte tgaaatgeag eteeetgtee 600
                                                                   634
aagtgccttg gagaactcac agcagcacgc ctta
<210> 541
<211> 221
<212> DNA
<213> Homo sapiens
<400> 541
 cacacaagca gcagagacca tgggaaccct ctcagcccct ccctgcacac agcgcatcaa 60
 atggaagggg ctcctgctca cagcatcact tttaaacttc tggaacctgc ccaccactgc 120
 ccaagtcacg attgaagccg agccaaccaa agtttccgag gggaaggatg ttcttctact 180
 tgtccacaat ttgccccaga atcttaccgg ctacatctgg t
```

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<211> 287
<212> DNA
<213> Homo sapiens
<400> 542
cctcttctac tatggcagga gatgtggcgt gctgttgcaa agttttcacg tcatcgtttc 60
ctggctagtt catttcatta agtggctaca tcctaacata tgcatttggt caaggttgca 120
gaagaggact gaagattgac tgccaagcta gtttgggtga agttcactcc agcaagtctc 180
aggccacaat ggggtggttt ggtttggttt ccttttaact ttccttttgt tatttgcttt 240
tctcctccac ctgtgtggta tattttttaa gcagaatttt attttt
<210> 543
<211> 274
<212> DNA
<213> Homo sapiens
<400> 543
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ggcacttaag cacaagcaga gtgcacagct gtccactggg ccattgtggt gtgagcttca 120
gatggtgaag catteteec agtgtatgte ttgtateega tatetaaege tttaaatgge 180
tactttggtt tctgtctgta agttaagacc ttggatgtgg tttaattgtt tgtcctcaaa 240
aggaataaaa cttttctgct gataagataa aaaa
<210> 544
<211> 307
<212> DNA
<213> Homo sapiens
<400> 544
ccaggtggtt gtcttattgc accatactcc ttgcttcctg atgctgggca atgaggcaga 60
tagcactggg tgtgagaatg atcaaggatc tggaccccaa agaatagact ggatggaaag 120
acaaactgca caggcagatg tttgcctcat aatagtcgta agtggagtcc tggaatttgg 180
acaagtgctg ttgggatata gtcaacttat tctttgagta atgtgactaa aggaaaaaac 240
tttgactttg cccaggcatg aaattcttcc taatgtcaga acagagtgca acccagtcac 300
actgtgg
<210> 545
<211> 570
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(570)
<223> n = A, T, C or G
<400> 545
accttagaaa tttgcaacca cctccctgaa agtcttctcc cacgttatta agtgcaatgt 60
ttatggtaaa tgtagaagca tcatgatgag gacgaagaga acgctgtcgt tcaggggagt 120
attttactac aaaattcagt agtgcaaatc ccttcgtata atagcctgca aagaccttca 180
gtgtaactgg ngcaatgaac tcccggataa aatgaagcca tacattctcc agatcaactt 240
gcttcatgtg gatatcatca gttgggacat tttcataacc accagatata cggctatcat 300
gatgttttcc cccagaccat ttgccgtaat, gttccatttc ttctaccaat tcatcacagg 360
ctttttcaga aaatatgggg aaccaaaaga catctggaca gggctgttca actatatttt 420
```

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cagtgaaaat ctttgaataa tcacggttta tatacttttc cttccagtcc acaggatttt 480
caaaaatctg ccagaggtca ttgttataat gggaagtatt gtaattagca gtggataata 540
gccttccaaa ttcatgtcta ttagaaatgt
<210> 546
<211> 589
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(589)
<223> n = A, T, C or G
<400> 546
aaaaatactt tttaccaaag gtgctatttc tctgtaaaac acttttttt ggcaagttga 60
ctttattctt caattattat cattatatta ttgtttttta atattttatt ttcttgacta 120
ggtattaagc ttttgtaatt atttttcagt agtcccacca cttcataggt ggaaggagtt 180
tggggttctt cctggtgcag gggctgaaat aacccagatg cccccaccct gccacatact 240
agatgcagcc catagttggc ccccctagct tccagcagtc cactatctgc cagaggagca 300
agggtgcctt agaccgaagc caggggaaga agcatcttca taaaaaactt tcaagatcca 360
aacattaatt tgtttttatt tattctgaga agttgaggca aatcagtatt cccaaggatg 420
gcgacaaggg cagccaagca gggcttagga tatcccagcc taccaatatg ctcattcgac 480
taactaggag ggtgagttgg coctgtctct tottttttct ggacctcagt ttocttcagt 540
                                                                   589
ggagcttggt aaaaatgcac taccntttga tttgataagg tataaatct
<210> 547
<211> 293
<212> DNA
<213> Homo sapiens
<400> 547
actcctatta ttgactgtag tcaatcaaac ataaaaaggt gaaagtaaaa tttaattttt 60
taccettatt ttactgacca atatggaagt tettggtate tttaaggetg acetteetgg 120
tattgtgtaa tgattgaatg tatctaaact gtaataattt gaaactgaca aacataacct 180
tctcagactt acaaaactat gttctttcta aagatacaga tttttattat tttattttga 240
ctaggaagga tttataaata aatgtaatga aaaatctttg atcttaataa agt
<210> 548
<211> 98
<212> DNA
<213> Homo sapiens
<400> 548
aaacaaaggt tgagatgtaa aaggtattaa attgatgttg ctggactgtc atagaaatta 60
cacccaaaga ggtatttatc tttacttttt tttgtaca
<210> 549
<211> 121
<212> DNA
<213> Homo sapiens
 <400> 549
acatgcatat ttcaaagacc tgttaatggc gtccactttg gattcttaca tgaaacgatt 60
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cagtgcacat tgtaagccta aggaccacgc aaaagggttt cccacatatt aagtattcag 120
                                                                   121
<210> 550
<211> 509
<212> DNA
<213> Homo sapiens
<400> 550
acaatagtat acattttata atgatgaact tataatgatt aagggacatt tctataaaaa 60
tactacaata gttttatgca caacttccca ttaaaaatga gatttcttat ttgtttgtct 120
gtttttactc tgggagtaat actttttaaa ttacctttac atatatagtc actggcatac 180
tgagaatata caatgatcct ggaaattgca gtaacaaaag cacacaacga ttatagtaac 240
tataagatac aataaaacaa ataaatgtga aagtagattc atgaaaatgt attcctttaa 300
aatattgttt tcctacaggc ctatttaaca agatgtttca ttttactgta tattttgtag 360
ttaatataaa tgttgctcta atcagattgc ttaaaagcat ttttattata tttatgttgt 420
tgaactaata tatgaaataa gtaaatgtag ctcccacaag gtaaacttca ttggtaagat 480
tgcactgttc tgattatgta agcatttgt
<210> 551
<211> 427
<212> DNA
<213> Homo sapiens
<400> 551
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aaaagaagct taagttttat catccttttt tttctcgtga attcttaaag gattatgctt 120
taatgctgtt atctatctta ttgttcttga aaatacctgc attttttggt atcatgttca 180
accaacatca ttatgaaatt aattagattc ccatggccat aaaatggctt taaagaatat 240
atatatattt ttaaagtagc ttgagaagca aattggcagg taatatttca tacctaaatt 300
aagactctga cttggattgt gaattataat gatatgcccc ttttcttata aaaacaaaaa 360
aaaaaataat gaaacacagt gaatttgtag agtgggggta tttgacatat tttacagggt 420
                                                                   427
ggagtgc
<210> 552
<211> 340
<212> DNA
<213> Homo sapiens
<400> 552
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ctgtcaagag aaatggtcca ccgtgtgtgt ggaatgcagc catcacacat tagtttctga 120
gattgcttct gtcttggttt tatggggaga tatttccatt tctagcatag gcttcaaggc 180
gctctaaata tccgcttgga aatactacaa aaacagtgtt tcaaaactgc tgtatccaaa 240
ggaaggtgcc actcgctgag ttgaatgcac acatcacaag gaagtttctg agaattcttc 300
                                                                   340
tgtctagatt catacgaaga aatcccgttt ccaacgaagg
<210> 553
<211> 549
<212> DNA
<213> Homo sapiens
<400> 553
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```

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ttcagacccg gggtggccca gaaaagcatg aagtaactgg ctgggtgctg gtatctcctc 180
taagtaagga agatgctgga gaatatgagt gccatgcatc caattcccaa ggacaggctt 240
cagcatcagc aaaaattaca gtggttgatg ccttacatga aataccagtg aaaaaaggtg 300
aaggtgccga gctataaacc tccagaatat tattagtctg catggttaaa agtagtcatg 360
tagttatatt cactggtttt acacagagaa atacaaaata aagatcacac atcaagacta 480
tctacaaaaa tttattatat atttacagaa gaaaagcatg catatcatta aacaaataaa 540
atacttttt
<210> 554
<211> 321
<212> DNA
<213> Homo sapiens
<400> 554
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actgaaaatg aagttaccat tootaggoca aatttttaga caaagcttto taaaaccato 120
tttataaagt aaattcagat atgcttacaa taaaaagaca taaaagattc atcctgagat 180
gaattotgag toaataacta aaaaccattt otaccagtgo atcactacca tgtaatccat 240
tctacgcaag ctctacaaat attgagtcaa atcctgtctg tcagaaaatg aagacccaat 300
                                                                321
aagtttgccg aagtattcag t
<210> 555
<211> 322
<212> DNA
<213> Homo sapiens
<400> 555
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gtactttgaa gttataacta atctgcctga agacttctca tgatggaaaa tcagccaagg 120
actaagcttc catagaaata cactttgtat ctggacctca aaattatggg aacatttact 180
taaacggatg atcatagctg aaaataatga tactgtcaat ttgagatagc agaagtttca 240
cacatcaaag taaaagattt gcatatcatt atactaaatg caaatgagtc gcttaaccct 300
tgacaaggtc aaagaaaact tt
<210> 556
<211> 286
<212> DNA
<213> Homo sapiens
<400> 556
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cgggccctcc tcttcaggaa tcttcctgaa gacatggccc agtcgaaggc ccaggatggc 120
ttttgctgcg gccccgtggg gtaggaggga cagagagaca gggagagtca gcctccacat 180
tcagaggcat cacaagtaat ggcacaattc ttcggatgac tgcagaaaat agtgttttgt 240
agttcaacaa ctcaagacga agcttatttc tgaggataag ctcttt
<210> 557
<211> 459
<212> DNA
<213> Homo sapiens
<400> 557
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<400> 560

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acagaagatg aataataatg aaaaactgtg attttttgac tatcacatac attgtgttaa 60
aaaacaggta aatataatga ctattactgt taagaaagac aaggaggaaa actgtttcaa 120
tgttcaggtt taaatactaa gcacaaaaat ataacaaatt ctgtgtctac aataattttt 180
gaagtgtata caagtgcatt gcaaatgagc tetttaaaat ttaaagteca ttteecettt 240
agccaagcat atgtctacat ttatgatttc tttctcttat tttaaagtct cttctggttt 300
agtttttaa aaagtttcat catggctgtc atcttggaat ctagcctcca gctcaaagct 360
gagacttcac gcatacatat tctcctttct gggtgcatct tcacctagtt tctccaagta 420
ttcagagtta aatagcacaa cttcttttat atgttccct
<210> 558
<211> 303
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(303)
<223> n = A, T, C or G
<400> 558
aaaaaataaa aaacaagaca acaatttagt agaagtaccn ctgggaggga ggggagggga 60
aaaaaggata tacaggggca ggngtattct ctgtacagag gtgcananaa aatttcacat 120
anctttanag aatgccttgt ggaaaaaaaa aaataggccc caatacttgt tactgccctt 180
tatcaaaact gtgtgcatga cctgcacaaa taaaatcaca aaacagtgtt gccacattct 240
tcaaggaaac aaagcaaaat ttagggggnt tcttttccct ctccttgtta aaagtcattt 300
                                                                   303
ttt
<210> 559
<211> 232
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(232)
<223> n = A, T, C or G
<400> 559
aaagcattta ttaagaattt actcaggcat gatggcccat acttgtaatc ccagctattg 60
ggaaggatga gatgggagga tggcttgagg ccagaggttt gagaccgacc agccagggca 120
acacagtgag accccttctc aaaaaaaaaa aaaaaaaaag agagagtgtg tgattagaag 180
                                                                    232
ctaaatagga aagttttgag cttcaagtca gngaggagta aaaaagattt tt
<210> 560
<211> 336
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(336)
<223> n = A, T, C or G
```

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ctctgcaaaa ataannataa aaaaataaat aaaattttaa aaataataaa attcactata 60
tacacatata aagaaataaa aagaagtete agttgeaget atttgteaaa attaatatee 120
atttcttttt atatacggtg aatattgcgc aattatagat ctggattttg aaccacttaa 180
tgaagcggca acaccaggtg ttttgaggtg ttggcattct tcgctgattt ggctgttccc 240
aatgtttaca ttatttaatc ttgcaaaaat ggttctgtgc acttggatgn gaaatgctgn 300
ccagntttat tttttttatg ttgntatcct tggatg
<210> 561
<211> 636
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(636)
<223> n = A, T, C or G
<400> 561
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aaatcaaata gaatcttata tctgtatgtt aaaatagagc acttacctga agtcagtggc 120
ctggatcata gccctggatc atttcccagt ctgtcctgtg ctgtgtgacc ttggacaagg 180
cgcttcatct ctctgggcct ctatttctcc atttgtaaaa caagtggctg cagtagatga 240
tggctgagag cccttcctgt tcccagatgc cttggtccaa agaccccacc cctctgctgg 300
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tatggacttg gttactgtaa taatgtcttg tttttagcca tgtaactaca aacagatatt 480
ctcttgatgt cttagtaaat ttgcatttga tatatcattg atgagatttt gttgttatgt 540
aatattettt ggetacgeat etgteeagea tettattaac cataataetg ngateattat 600
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<210> 562
<211> 708
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(708)
<223> n = A, T, C or G
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tggagaacaa ctaaggggca ccaaaccctc tgaggtttta ctttaaggtt cgctgtatgt 180
ttgccttgga caaaaaggct acctaccacg tgctatccag taatatactt aaataagcca 240
atacttagat ctactgtaag gcagatgcta attataaggc attaagtaag caaatagtgc 300
cctcagctac tgcagaagaa aagtcccact gaggaaaaga aagtcttgtg atttttaaag 360
gcaagttttc aagtgctctc atagttctat cctctaattc cattaaatcc atactaggag 420
cgtcagtgag ggttttcata gcttttggaa atactttggt ctctgaactg taattagcaa 480
gaagtaaaaa cagaaacgtc aaacgtcaaa tgtttgcttt gttacctgga ggactaaatg 540
tagatgtctt tagtatactt tgtatgttct taatattgga agataatttt gtgaatctgt 600
 agattttatt ttttcagtct taccttacaa atttcttttc tatgaataat agaggactta 660
                                                                    708
 cngcactctg ccatttgtta atgaaaggaa ggcngangat ttagaaag
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<211> 290
<212> DNA
<213> Homo sapiens
<400> 563
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ctgtataata ggcttgagag atatccttca tttgcctgct tgtcctggta gttaagattt 180
caatcaaggc atcttcgttt gttcccgcgc ccttcatgga tttctttagc tgctttgcat 240
caaagactgc tggtggagtc actagggcca ccatgagatg ctcaaagtgg
<210> 564
<211> 530
<212> DNA
<213> Homo sapiens
<400> 564
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accatacgct ccaaaagatg gctgtgatag atcttgtgaa gcaattactg agcagatcaa 120
gatctttggg aaggaacact aaagatgttt tgaatgaatt atagtccact ggcattttag 180
tgtatttttt tttctttta gaaacacaca tttctaaaaa tgtcatgtta cattcctgca 240
tgtccctttt gatagcatta gtggatccat tggatttctt ttttcttttt gtgagacagc 300
ttttagtctt acctgaattt atgtgtgttt ttccgacagt ggttaataat tatattggtg 360
atgtagcagc aattgtgttg gcagggtttt catatattat tagtaattaa cactaactgt 420
tggactgact tgtgtcgata gcgctcacgc aagcatggtt aacgtcccta aaacccgccg 480
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gactttctgt aagaagtgtg gcaagcacca accccataaa gtgacacagt
<210> 565
<211> 450
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(450)
<223> n = A, T, C or G
<400> 565
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gtgtgttaag tttttcatct gtgcatcaaa tcacaaaaag aataaataga gctttttcct 120
ttatcagtcc cttgggcaca gcaggtcctg aacaccctgc tctacaatgt tgcatcaaga 180
gttcaaacaa caaaataaaa aatattaaga ggaaatcccc atcctgtgac ttgagtccct 240
taagtctaca ggggctggtg acctcttttt gctaatagga aaatcacatt actacaaaat 300
ggggagaaaa ctgtttgcct gtggtagaca cctgcacgca taggattgaa gacagtacag 360
gctgctgtac agagaagcgc ctctcacatc tgaactgcat actgagcggg caagtcggtt 420
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<210> 566
<211> 563
<212> DNA
<213> Homo sapiens
<400> 566
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<212> DNA

<213> Homo sapiens

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ttcagacccg gggtggccca gaaaagcatg aagtaactgg ctgggtgctg gtatctcctc 180
taagtaagga agatgctgga gaatatgagt gccatgcatc caattcccaa ggacaggctt 240
cagcatcagc aaaaattaca gtggttgatg ccttacatga aataccagtg aaaaaaggtg 300
aaggtgccga gctataaacc tccagaatat tattagtctg catggttaaa agtagtcatg 360
tagttatatt cactggtttt acacagagaa atacaaaata aagatcacac atcaagacta 480
tctacaaaaa tttattatat atttacagaa gaaaagcatg catatcatta aacaaataaa 540
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atactttta tcacaaaaaa aaa
<210> 567
<211> 424
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(424)
<223> n = A, T, C or G
<400> 567
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gaataccett etecatagea ggtgeaatge tgaetgetea aggegtgegt gegegegeae 120
acacacaca acacacaca atacatacte teacacaene atettteeaa ttaaaetgea 180
ggtagaatga gattttgtgt tattcaaaaa atttgtaagt gatcaaaanc actgctatgg 240
aatgcctgtt tatctgcctt tgntctggtt aaaatctcat aaaaatacat tcaacaggaa 300
aacatanatt gtatgtgtat aaatatatat gtatatatat atattatata cacatgcaca 360
caaatacttt tgttttttga agcataagat agttacataa atactcctat aattgctaaa 420
                                                                 424
gttt
<210> 568
<211> 392
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(392)
<223> n = A, T, C or G
<400> 568
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tttgaagaag aaaggatggg aggtggtgga gtcggacctc tatgccatga acttcaatcc 120
catcatttcc agaaaggaca tcacaggtaa actgaaggac cctgcgaact ttcagtatcc 180
tgccgagtct gttctggctt ataaagaagg ccatctgagc ccagatattg tgggttganc 240
aaaagaaagc ttggaagccn caagaacctt gtgatattcc agttccccct gcantgggtt 300
tgggaagtcc ctgccntttt gaaagctggt ttgaagcgaa tgttcatagg aaagtttgct 360
taccacttac cctgcccatg gtangacaaa ag
<210> 569
<211> 559
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<210> 571 <211> 261 <212> DNA <213> Homo	sapiens					
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<210> 573
<211> 619
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(619)
<223> n = A, T, C or G
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gaattgaaac cccagaagat aactacaaca aaaacatgtt aattttttt taaaaatgat 120
gattcaaagg cagatttgaa gggaagtaat atttaggtgg cagaagaagg caaatgcagc 180
ctctgaaggg aactgttcta attattacct aaaaaataaa gttacacaac tatattcaag 240
gacatgagat aaagcactgc ttgaaaacca gaatgactga acagttaggt gaaaaggaac 300
agctgaaata ggaaggggaa atggactgaa gaataatttg aatcgggaca gtgatccatc 360
agtectagat gettetggta tgtaaatate ttgaateaca ttgttteett tettetgaaa 420
tetcaaagga gaattetcae ageactaeat taaggttgee attttgttag gatteaaaat 480
ttcaatccag tagccatcag gatcttgaat aaatgccagg cctttcattt taccatcatc 540
aggtttcttc acaaatttga ctccagtctt caaccttttc aagcctgatc atcaggaaca 600
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caattccata tgaccgatc
<210> 574
<211> 202
<212> DNA
<213> Homo sapiens
<400> 574
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teetgagaag geetgteagt tggacagteg etattggaga ataacaaatg etaagggtga 120
cgtggaagaa gttcaaggac ctggagtagt tggtgaattt ccaatcatca gcccaggtcg 180
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 ggtatatgaa tacacaagct gt
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 <211> 311
 <212> DNA
 <213> Homo sapiens
 <400> 575
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 gtgtgaatta taaagaaaag catgagaatg actctaagtt caacaaacat gggtgaatct 180
 ctatgtgctc ccagtgtcct ggatgggctc cccagcaagc cattcctcct tcctgttctg 240
 atattactat tettettae attgegetaa ggaggacaaa agatgagaga tgaaaataaa 300
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 gctttgcctt t
 <210> 576
 <211> 134
 <212> DNA
 <213> Homo sapiens
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<221> misc feature
<222> (1)...(134)
<223> n = A, T, C or G
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cttcaaaagt cgct
<210> 577
<211> 488
<212> DNA
<213> Homo sapiens
<400> 577
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agttgctgca aacctgaccc ctgctcagta aagcacttgc aaccgtctgt tatgctgtga 120
cacatggccc ctcccctgc caggagcttt ggacctaatc caagcatccc tttgcccaga 180
aagaagatgg gggaggaggc agtaataaaa agattgaagt attttgctgg aataagttca 240
aattettetg aacteaaact gaggaattte acetgtaaac etgagtegta cagaaagetg 300
cctggtatat ccaaaagctt tttattcctc ctgctcatat tgtgattctg cctttgggga 360
cttttcttaa accttcagtt atgattttt tttcatacac ttattggaac tctgcttgat 420
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                                                                   488
ttgcattt
<210> 578
<211> 476
<212> DNA
<213> Homo sapiens
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catcaacttg cagaaagaaa tataaatgac atttcaagga tagaagtata cctgattttt 180
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atgtctcttc tgagctattt catctatttt tggcagtctg aatttttaaa acccatttaa 360
atttttttcc ttaccttttt atttgcatgt ggatcaacca tcgctttatt ggctgagata 420
tgaacatatt gttgaaaggt aatttgagag aaatatgaag aactgaggaa aaaaaa
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<211> 246
<212> DNA
<213> Homo sapiens
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ctagaatgaa gccaccaatt tcaatgtgac caggcaatgg cagccagcac tgccttacac 120
tggtttgatt ctgattccct aattctggcc actgcaggtg atgagtaagg gtggggatca 180
gggaggaagt ccagaagcca gtctttgtct ccctttcctg cttatattta agtgcctatt 240
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tacatg
<210> 580
<211> 615
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<212> DNA
<213> Homo sapiens
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<221> misc feature
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ccaggtgtgg tectetagae actggeteeg attgetgeee ttgaggatgt agtggteatt 180
gcacataaac gtgattttgt cacttacatt cacaggccct gaagaactga actctccatt 240
caccagcaca ggatcaggac agtggcccaa gcggcactca gtagtggtgt tatcccactc 300
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qtaagtcccc agaatctgtc cttccacctc ctttgcgaca aatatgctat tgtccactgg 420
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cgcacaccaa aaaaacatct ggtgatcaaa gtcctctccc caggctggaa ttcacccagc 540
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ttaactgact tccaa
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<211> 576
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qaqttctqqt caaagaaaga aagtttagaa gctgagacac aaagggttgg gagctgatga 180
aactcacaaa tgatggtagg aagaagctct cgacaatacc cgttggcaag gagtctgcct 240
ccatgctgca gtgttcgagt ggattgtagg tgcaagatgg aaaggattgt aggtgcaagc 300
tqtccaqaqa aaaqaqtcct tgttccagcc ctattctgcc actcctgaca gggtgacctt 360
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<212> DNA
<213> Homo sapiens
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cagtgtgccc aggatatgaa ccatgaatac atgtcctggt atcgacaaga cccaggcatg 180
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gctccctccc agacatctgt gtacttctgt gccagcagtt actcagtcgg ggagggcggg 360
gattcacccc tccactttgg gaatgggacc aggctcactg tgacagagga cctgaacaag 420
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Ser Glu Glu 35		Leu Th	r Ile . 40	Asn Cy	s Thr	Tyr	Thr 45	Ala	Thr	Gly	
Tyr Pro Ser 50	Leu Phe	Trp Ty 5	_	Gln Ty	r Pro	Gly 60	Glu	Gly	Leu	Gln	
Leu Leu Leu 65	ı Lys Ala	Thr Ly 70	s Ala	Asp As <sub>l</sub>	Lys 75	Gly	Ser	Asn	Lys	Gly 80	
Phe Glu Ala	Thr Tyr 85	Arg Ly	s Glu	Thr Th	_	Phe	His	Leu	Glu 95	Lys	

Gly Ser Val Gln Val Ser Asp Ser Ala Val Tyr Phe Cys Ala Pro Asn 105 Pro Ser Leu Gln Gly Gly Ser Glu Lys Leu Val Phe Gly Lys Gly Thr 115 Lys Leu Thr Val Asn Pro Tyr Ile Gln Asn Pro Asp Pro Ala Val Tyr Gln Leu Arg Asp Ser Lys Ser Ser Asp Lys Ser Val Cys Leu Phe Thr Asp Phe Asp Ser Gln Thr Asn Val Ser Gln Ser Lys Asp Ser Asp Val 165 Tyr Ile Thr Asp Lys Thr Val Leu Asp Met Arg Ser Met Asp Phe Lys 185 Ser Asn Ser Ala Val Ala Trp Ser Asn Lys Ser Asp Phe Ala Cys Ala 195 200 Asn Ala Phe Asn Asn Ser Ile Ile Pro Glu Asp Thr Phe Phe Pro Ser 215 Pro Glu Ser Ser Cys Asp Val Lys Leu Val Glu Lys Ser Phe Glu Thr 235 230 Asp Thr Asn Leu Asn Phe Gln Asn Leu Ser Val Ile Gly Phe Arg Ile 245 Leu Leu Lys Val Ala Gly Phe Asn Leu Leu Met Thr Leu Arg Leu 265 Trp Ser Ser 275 <210> 585 <211> 312 <212> PRT <213> Homo sapiens <400> 585 Met Ser Ile Gly Leu Leu Cys Cys Ala Ala Leu Ser Leu Leu Trp Ala Gly Pro Val Asn Ala Gly Val Thr Gln Thr Pro Lys Phe Gln Val Leu

Lys Thr Gly Gln Ser Met Thr Leu Gln Cys Ala Gln Asp Met Asn His

Glu Tyr Met Ser Trp Tyr Arg Gln Asp Pro Gly Met Gly Leu Arg Leu

40

35

-

266

50 55 60 Ile His Tyr Ser Val Gly Ala Gly Ile Thr Asp Gln Gly Glu Val Pro 70 75 Asn Gly Tyr Asn Val Ser Arg Ser Thr Thr Glu Asp Phe Pro Leu Arg 90 Leu Leu Ser Ala Ala Pro Ser Gln Thr Ser Val Tyr Phe Cys Ala Ser 105 Ser Tyr Ser Val Gly Gly Gly Asp Ser Pro Leu His Phe Gly Asn Gly Thr Arg Leu Thr Val Thr Glu Asp Leu Asn Lys Val Phe Pro Pro 135 Glu Val Ala Val Phe Glu Pro Ser Glu Ala Glu Ile Ser His Thr Gln 145 150 155 160 Lys Ala Thr Leu Val Cys Leu Ala Thr Gly Phe Phe Pro Asp His Val 165 170 Glu Leu Ser Trp Trp Val Asn Gly Lys Glu Val His Ser Gly Val Ser 185 Thr Asp Pro Gln Pro Leu Lys Glu Gln Pro Ala Leu Asn Asp Ser Arg 195 Tyr Cys Leu Ser Ser Arg Leu Arg Val Ser Ala Thr Phe Trp Gln Asn 215 Pro Arg Asn His Phe Arg Cys Gln Val Gln Phe Tyr Gly Leu Ser Glu 225 230 235 240 Asn Asp Glu Trp Thr Gln Asp Arg Ala Lys Pro Val Thr Gln Ile Val Ser Ala Glu Ala Trp Gly Arg Ala Asp Cys Gly Phe Thr Ser Val Ser Tyr Gln Gln Gly Val Leu Ser Ala Thr Ile Leu Tyr Glu Ile Leu Leu 280 275 285 Gly Lys Ala Thr Leu Tyr Ala Val Leu Val Ser Ala Leu Val Leu Met 295 300 Ala Met Val Lys Arg Lys Asp Phe 305

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<213> Homo sapiens
<400> 586
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Leu Ser Gln Thr Glu Leu Arg Lys Lys Glu Arg Lys Lys Arg Glu
Arg Lys Phe Gln Ala Asn Cys Gly Ile Asp Phe Ile Ile Phe Trp Ile
Phe Trp Ile Leu Leu Phe Ser His His Trp Ile Gln Glu Ser Leu Leu
                         55
Cys Pro Pro Ser Pro Lys Glu Val Thr Cys Arg Glu Met Leu Thr Gly
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Gly Cys Leu Pro Trp Ala Thr Arg Ser His Leu Gly Arg Arg Lys Cys
Ser
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gactgccagg caataatgaa ggttctttta ctgaaggatg cgaaggaaga tgactgtggc 180
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<212> DNA
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gccccagttc agggcagcat ccatagccca caagccagcg tgggtgggc gggggtggtc 240
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